



# SEQUENCE LISTING

<110> Srivastava, Pramod

<120> COMPLEXES OF ALPHA (2) MACROGLOBULIN AND ANTIGENIC MOLECULES FOR IMMUNOTHERAPY

<130> 8449-178-999

<140> 09/873,403

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<151> 2000-06-02

<160> 57

<170> PatentIn version 3.2

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Gly His Met Glu Val Leu Arg Gly His Glu Phe Leu Ser His Pro	
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ttt gcc gtg aca ctg tac ggt ggg gag gtg tac tgg acc gac tgg	4932
Phe Ala Val Thr Leu Tyr Gly Gly Glu Val Tyr Trp Thr Asp Trp	
1485 1490 1495	
cga aca aat aca ctg gct aag gcc aac aag tgg act ggc cac aac	4977
Arg Thr Asn Thr Leu Ala Lys Ala Asn Lys Trp Thr Gly His Asn	
1500 1505 1510	
gtc acc gtg gta cag agg acc aac acc cag ccc ttc gac ctg cag	5022
Val Thr Val Val Gln Arg Thr Asn Thr Gln Pro Phe Asp Leu Gln	
1515 1520 1525	
gtg tat cac cct tcc cgg cag ccc atg gct cca aac cca tgt gag	5067
Val Tyr His Pro Ser Arg Gln Pro Met Ala Pro Asn Pro Cys Glu	
1530 1535 1540	
gcc aat ggc ggc cgg ggc ccc tgt tcc cat ctg tgc ctc atc aac	5112
Ala Asn Gly Gly Arg Gly Pro Cys Ser His Leu Cys Leu Ile Asn	
1545 1550 1555	
tac aac cgg acc gtc tcc tgg gcc tgt ccc cac ctc atg aag ctg	5157
Tyr Asn Arg Thr Val Ser Trp Ala Cys Pro His Leu Met Lys Leu	
1560 1565 1570	
cac aag gac aac acc acc tgc tat gag ttt aag aag ttc ctg ctg	5202
His Lys Asp Asn Thr Thr Cys Tyr Glu Phe Lys Lys Phe Leu Leu	
1575 1580 1585	
tac gca cgt cag atg gag atc cgg ggc gtg gac ctg gat gcc ccg	5247
Tyr Ala Arg Gln Met Glu Ile Arg Gly Val Asp Leu Asp Ala Pro	
1590 1595 1600	
tac tac aat tat atc atc tcc ttc acg gtg cct gat atc gac aat	5292
Tyr Tyr Asn Tyr Ile Ile Ser Phe Thr Val Pro Asp Ile Asp Asn	
1605 1610 1615	
gtc acg gtg ctg gac tat gat gcc cga gag cag cga gtt tac tgg	5337
Val Thr Val Leu Asp Tyr Asp Ala Arg Glu Gln Arg Val Tyr Trp	
1620 1625 1630	
tct gat gtg cgg act caa gcc atc aaa agg gca ttt atc aac ggc	5382
Ser Asp Val Arg Thr Gln Ala Ile Lys Arg Ala Phe Ile Asn Gly	
1635 1640 1645	
act ggc gtg gag acc gtt gtc tct gca gac ttg ccc aac gcc cac	5427
Thr Gly Val Glu Thr Val Val Ser Ala Asp Leu Pro Asn Ala His	
1650 1655 1660	
ggg ctg gct gtg gac tgg gtc tcc cga aat ctg ttt tgg aca agt	5472
Gly Leu Ala Val Asp Trp Val Ser Arg Asn Leu Phe Trp Thr Ser	
1665 1670 1675	
tac gac acc aac aag aag cag att aac gtg gcc cgg ctg gac ggc	5517
Tyr Asp Thr Asn Lys Lys Gln Ile Asn Val Ala Arg Leu Asp Gly	
1680 1685 1690	
tcc ttc aag aat gcg gtg gtg cag ggc ctg gag cag ccc cac ggc	5562
Ser Phe Lys Asn Ala Val Val Gln Gly Leu Glu Gln Pro His Gly	
1695 1700 1705	
ctg gtc gtc cac ccg ctt cgt ggc aag ctc tac tgg act gat ggg	5607



Leu	Val	Val	His	Pro	Leu	Arg	Gly	Lys	Leu	Tyr	Trp	Thr	Asp	Gly		
		1710					1715					1720				
gac	aac	atc	agc	atg	gcc	aac	atg	gat	ggg	agc	aac	cac	act	ctg		5652
Asp	Asn	Ile	Ser	Met	Ala	Asn	Met	Asp	Gly	Ser	Asn	His	Thr	Leu		
		1725					1730					1735				
ctc	ttc	agt	ggc	cag	aag	ggc	cct	gtg	ggg	ttg	gcc	att	gac	ttc		5697
Leu	Phe	Ser	Gly	Gln	Lys	Gly	Pro	Val	Gly	Leu	Ala	Ile	Asp	Phe		
		1740					1745					1750				
cct	gag	agc	aaa	ctc	tac	tgg	atc	agc	tct	ggg	aac	cac	aca	atc		5742
Pro	Glu	Ser	Lys	Leu	Tyr	Trp	Ile	Ser	Ser	Gly	Asn	His	Thr	Ile		
		1755					1760					1765				
aac	cgt	tgc	aat	ctg	gat	ggg	agc	gag	ctg	gag	gtc	atc	gac	acc		5787
Asn	Arg	Cys	Asn	Leu	Asp	Gly	Ser	Glu	Leu	Glu	Val	Ile	Asp	Thr		
		1770					1775					1780				
atg	cgg	agc	cag	ctg	ggc	aag	gcc	act	gcc	ctg	gcc	atc	atg	ggg		5832
Met	Arg	Ser	Gln	Leu	Gly	Lys	Ala	Thr	Ala	Leu	Ala	Ile	Met	Gly		
		1785					1790					1795				
gac	aag	ctg	tgg	tgg	gca	gat	cag	gtg	tca	gag	aag	atg	ggc	acg		5877
Asp	Lys	Leu	Trp	Trp	Ala	Asp	Gln	Val	Ser	Glu	Lys	Met	Gly	Thr		
		1800					1805					1810				
tgc	aac	aaa	gcc	gat	ggc	tct	ggg	tcc	gtg	gtg	ctg	cgg	aac	agt		5922
Cys	Asn	Lys	Ala	Asp	Gly	Ser	Gly	Ser	Val	Val	Leu	Arg	Asn	Ser		
		1815					1820					1825				
acc	acg	ttg	gtt	atg	cac	atg	aag	gtg	tat	gac	gag	agc	atc	cag		5967
Thr	Thr	Leu	Val	Met	His	Met	Lys	Val	Tyr	Asp	Glu	Ser	Ile	Gln		
		1830					1835					1840				
cta	gag	cat	gag	ggc	acc	aac	ccc	tgc	agt	gtc	aac	aac	gga	gac		6012
Leu	Glu	His	Glu	Gly	Thr	Asn	Pro	Cys	Ser	Val	Asn	Asn	Gly	Asp		
		1845					1850					1855				
tgt	tcc	cag	ctc	tgc	ctg	cca	aca	tca	gag	acg	act	cgc	tcc	tgt		6057
Cys	Ser	Gln	Leu	Cys	Leu	Pro	Thr	Ser	Glu	Thr	Thr	Arg	Ser	Cys		
		1860					1865					1870				
atg	tgt	aca	gcc	ggt	tac	agc	ctc	cgg	agc	gga	cag	cag	gcc	tgt		6102
Met	Cys	Thr	Ala	Gly	Tyr	Ser	Leu	Arg	Ser	Gly	Gln	Gln	Ala	Cys		
		1875					1880					1885				
gag	ggt	gtg	ggc	tct	ttt	ctc	ctg	tac	tct	gta	cat	gag	gga	att		6147
Glu	Gly	Val	Gly	Ser	Phe	Leu	Leu	Tyr	Ser	Val	His	Glu	Gly	Ile		
		1890					1895					1900				
cgg	ggg	att	cca	cta	gat	ccc	aat	gac	aag	tcg	gat	gcc	ctg	gtc		6192
Arg	Gly	Ile	Pro	Leu	Asp	Pro	Asn	Asp	Lys	Ser	Asp	Ala	Leu	Val		
		1905					1910					1915				
cca	gtg	tcc	gga	act	tca	ctg	gct	gtc	gga	atc	gac	ttc	cat	gcc		6237
Pro	Val	Ser	Gly	Thr	Ser	Leu	Ala	Val	Gly	Ile	Asp	Phe	His	Ala		
		1920					1925					1930				
gaa	aat	gac	act	att	tat	tgg	gtg	gat	atg	ggc	cta	agc	acc	atc		6282
Glu	Asn	Asp	Thr	Ile	Tyr	Trp	Val	Asp	Met	Gly	Leu	Ser	Thr	Ile		

1935				1940				1945							
agc	agg	gcc	aag	cgt	gac	cag	aca	tgg	cga	gag	gat	gtg	gtg	acc	6327
Ser	Arg	Ala	Lys	Arg	Asp	Gln	Thr	Trp	Arg	Glu	Asp	Val	Val	Thr	
1950				1955				1960							
aac	ggt	att	ggc	cgt	gtg	gag	ggc	atc	gcc	gtg	gac	tgg	atc	gca	6372
Asn	Gly	Ile	Gly	Arg	Val	Glu	Gly	Ile	Ala	Val	Asp	Trp	Ile	Ala	
1965				1970				1975							
ggc	aac	ata	tac	tgg	acg	gac	cag	ggc	ttc	gat	gtc	atc	gag	gtt	6417
Gly	Asn	Ile	Tyr	Trp	Thr	Asp	Gln	Gly	Phe	Asp	Val	Ile	Glu	Val	
1980				1985				1990							
gcc	cgg	ctc	aat	ggc	tct	ttt	cgt	tat	gtg	gtc	att	tcc	cag	ggg	6462
Ala	Arg	Leu	Asn	Gly	Ser	Phe	Arg	Tyr	Val	Val	Ile	Ser	Gln	Gly	
1995				2000				2005							
ctg	gac	aag	cct	cgg	gcc	atc	act	gtc	cac	cca	gag	aag	ggg	tac	6507
Leu	Asp	Lys	Pro	Arg	Ala	Ile	Thr	Val	His	Pro	Glu	Lys	Gly	Tyr	
2010				2015				2020							
ttg	ttc	tgg	acc	gag	tgg	ggg	cat	tac	cca	cgt	att	gag	cgg	tct	6552
Leu	Phe	Trp	Thr	Glu	Trp	Gly	His	Tyr	Pro	Arg	Ile	Glu	Arg	Ser	
2025				2030				2035							
cgc	ctt	gat	ggc	aca	gag	aga	gtg	gtg	ttg	gtt	aat	gtc	agc	atc	6597
Arg	Leu	Asp	Gly	Thr	Glu	Arg	Val	Val	Leu	Val	Asn	Val	Ser	Ile	
2040				2045				2050							
agc	tgg	ccc	aat	ggc	atc	tca	gta	gac	tat	cag	ggc	ggc	aag	ctc	6642
Ser	Trp	Pro	Asn	Gly	Ile	Ser	Val	Asp	Tyr	Gln	Gly	Gly	Lys	Leu	
2055				2060				2065							
tac	tgg	tgt	gat	gct	cgg	atg	gac	aag	atc	gag	cgc	atc	gac	ctg	6687
Tyr	Trp	Cys	Asp	Ala	Arg	Met	Asp	Lys	Ile	Glu	Arg	Ile	Asp	Leu	
2070				2075				2080							
gaa	acg	ggc	gag	aac	cgg	gag	gtg	gtc	ctg	tcc	agc	aat	aac	atg	6732
Glu	Thr	Gly	Glu	Asn	Arg	Glu	Val	Val	Leu	Ser	Ser	Asn	Asn	Met	
2085				2090				2095							
gat	atg	ttc	tcc	gtg	tcc	gtg	ttt	gag	gac	ttc	atc	tac	tgg	agt	6777
Asp	Met	Phe	Ser	Val	Ser	Val	Phe	Glu	Asp	Phe	Ile	Tyr	Trp	Ser	
2100				2105				2110							
gac	aga	act	cac	gcc	aat	ggc	tcc	atc	aag	cgc	ggc	tgc	aaa	gac	6822
Asp	Arg	Thr	His	Ala	Asn	Gly	Ser	Ile	Lys	Arg	Gly	Cys	Lys	Asp	
2115				2120				2125							
aat	gct	aca	gac	tcc	gtg	cct	ctg	agg	aca	ggc	att	ggg	gtt	cag	6867
Asn	Ala	Thr	Asp	Ser	Val	Pro	Leu	Arg	Thr	Gly	Ile	Gly	Val	Gln	
2130				2135				2140							
ctt	aaa	gac	atc	aag	gtc	ttc	aac	agg	gac	agg	cag	aag	ggg	acc	6912
Leu	Lys	Asp	Ile	Lys	Val	Phe	Asn	Arg	Asp	Arg	Gln	Lys	Gly	Thr	
2145				2150				2155							
aat	gtg	tgc	gcg	gta	gcc	aac	ggc	ggg	tgc	cag	cag	ctc	tgc	ttg	6957
Asn	Val	Cys	Ala	Val	Ala	Asn	Gly	Gly	Cys	Gln	Gln	Leu	Cys	Leu	
2160				2165				2170							

tat cgg ggt ggc gga cag cga gcc tgt gcc tgt gcc cac ggg atg	7002
Tyr Arg Gly 2175 Gly Gly Gln Arg Ala Cys Ala Cys Ala His Gly Met 2185 2185	
ctg gca gaa gac ggg gcc tca tgc cga gag tac gct ggc tac ctg	7047
Leu Ala Glu 2190 Asp Gly Ala Ser Cys Arg Glu Tyr Ala Gly Tyr Leu 2195 2200	
ctc tac tca gag cgg acc atc ctc aag agc atc cac ctg tcg gat	7092
Leu Tyr Ser 2205 Glu Arg Thr Ile Leu Lys Ser Ile His Leu Ser Asp 2210 2215	
gag cgt aac ctc aac gca ccg gtg cag ccc ttt gaa gac ccc gag	7137
Glu Arg Asn 2220 Leu Asn Ala Pro Val Gln Pro Phe Glu Asp Pro Glu 2225 2230	
cac atg aaa aat gtc atc gcc ctg gcc ttt gac tac cga gca ggc	7182
His Met Lys 2235 Asn Val Ile Ala Leu Ala Phe Asp Tyr Arg Ala Gly 2240 2245	
acc tcc ccg ggg acc cct aac cgc atc ttc ttc agt gac atc cac	7227
Thr Ser Pro 2250 Gly Thr Pro Asn Arg Ile Phe Phe Ser Asp Ile His 2255 2260	
ttt ggg aac atc cag cag atc aat gac gat ggc tcg ggc agg acc	7272
Phe Gly Asn 2265 Ile Gln Gln Ile Asn Asp Asp Gly Ser Gly Arg Thr 2270 2275	
acc atc gtg gaa aat gtg ggc tct gtg gaa ggc ctg gcc tat cac	7317
Thr Ile Val 2280 Glu Asn Val Gly Ser Val Glu Gly Leu Ala Tyr His 2285 2290	
cgt ggc tgg gac aca ctg tac tgg aca agc tac acc aca tcc acc	7362
Arg Gly Trp 2295 Asp Thr Leu Tyr Trp Thr Ser Tyr Thr Thr Ser Thr 2300 2305	
atc acc cgc cac acc gtg gac cag act cgc cca ggg gcc ttc gag	7407
Ile Thr Arg 2310 His Thr Val Asp Gln Thr Arg Pro Gly Ala Phe Glu 2315 2320	
agg gag aca gtc atc acc atg tcc gga gac gac cac ccg aga gcc	7452
Arg Glu Thr 2325 Val Ile Thr Met Ser Gly Asp Asp His Pro Arg Ala 2330 2335	
ttt gtg ctg gat gag tgc cag aac ctg atg ttc tgg acc aat tgg	7497
Phe Val Leu 2340 Asp Glu Cys Gln Asn Leu Met Phe Trp Thr Asn Trp 2345 2350	
aac gag ctc cat cca agc atc atg cgg gca gcc cta tcc gga gcc	7542
Asn Glu Leu 2355 His Pro Ser Ile Met Arg Ala Ala Leu Ser Gly Ala 2360 2365	
aac gtc ctg acc ctc att gag aag gac atc cgc acg ccc aat ggg	7587
Asn Val Leu 2370 Thr Leu Ile Glu Lys Asp Ile Arg Thr Pro Asn Gly 2375 2380	
ttg gcc atc gac cac cgg gcg gag aag ctg tac ttc tcg gat gcc	7632
Leu Ala Ile 2385 Asp His Arg Ala Glu Lys Leu Tyr Phe Ser Asp Ala 2390 2395	

acc ttg gac	aag atc gag cgc tgc	gag tac gac ggc tcc	cac cgc	7677
Thr Leu Asp	Lys Ile Glu Arg Cys	Glu Tyr Asp Gly Ser	His Arg	
2400	2405	2410		
tat gtg atc	cta aag tcg gag ccc	gtc cac ccc ttt ggg	ttg gcg	7722
Tyr Val Ile	Leu Lys Ser Glu Pro	Val His Pro Phe Gly	Leu Ala	
2415	2420	2425		
gtg tac gga	gag cac att ttc tgg	act gac tgg gtg cgg	cgg gct	7767
Val Tyr Gly	Glu His Ile Phe Trp	Thr Asp Trp Val Arg	Arg Ala	
2430	2435	2440		
gtg cag cga	gcc aac aag tat gtg	ggc agc gac atg aag	ctg ctt	7812
Val Gln Arg	Ala Asn Lys Tyr Val	Gly Ser Asp Met Lys	Leu Leu	
2445	2450	2455		
cgg gtg gac	att ccc cag caa ccc	atg ggc atc atc gcc	gtg gcc	7857
Arg Val Asp	Ile Pro Gln Gln Pro	Met Gly Ile Ile Ala	Val Ala	
2460	2465	2470		
aat gac acc	aac agc tgt gaa ctc	tcc ccc tgc cgt atc	aac aat	7902
Asn Asp Thr	Asn Ser Cys Glu Leu	Ser Pro Cys Arg Ile	Asn Asn	
2475	2480	2485		
gga ggc tgc	cag gat ctg tgt ctg	ctc acc cac caa ggc	cac gtc	7947
Gly Gly Cys	Gln Asp Leu Cys Leu	Leu Thr His Gln Gly	His Val	
2490	2495	2500		
aac tgt tcc	tgt cga ggg ggc cgg	atc ctc cag gag gac	ttc acc	7992
Asn Cys Ser	Cys Arg Gly Gly Arg	Ile Leu Gln Glu Asp	Phe Thr	
2505	2510	2515		
tgc cgg gct	gtg aac tcc tct tgt	cgg gca caa gat gag	ttt gag	8037
Cys Arg Ala	Val Asn Ser Ser Cys	Arg Ala Gln Asp Glu	Phe Glu	
2520	2525	2530		
tgt gcc aat	ggg gaa tgt atc agc	ttc agc ctc acc tgt	gat ggc	8082
Cys Ala Asn	Gly Glu Cys Ile Ser	Phe Ser Leu Thr Cys	Asp Gly	
2535	2540	2545		
gtc tcc cac	tgc aag gac aag tcc	gat gag aag ccc tcc	tac tgc	8127
Val Ser His	Cys Lys Asp Lys Ser	Asp Glu Lys Pro Ser	Tyr Cys	
2550	2555	2560		
aac tca cgc	cgc tgc aag aag act	ttc cgc cag tgt aac	aat ggc	8172
Asn Ser Arg	Arg Cys Lys Lys Thr	Phe Arg Gln Cys Asn	Asn Gly	
2565	2570	2575		
cgc tgt gta	tcc aac atg ctg tgg	tgc aat ggg gtg gat	tac tgt	8217
Arg Cys Val	Ser Asn Met Leu Trp	Cys Asn Gly Val Asp	Tyr Cys	
2580	2585	2590		
ggg gat ggc	tct gat gag ata cct	tgc aac aag act gcc	tgt ggt	8262
Gly Asp Gly	Ser Asp Glu Ile Pro	Cys Asn Lys Thr Ala	Cys Gly	
2595	2600	2605		
gtg ggt gag	ttc cgc tgc cgg gat	ggg tcc tgc atc ggg	aac tcc	8307
Val Gly Glu	Phe Arg Cys Arg Asp	Gly Ser Cys Ile Gly	Asn Ser	
2610	2615	2620		
agt cgc tgc	aac cag ttt gtg gat	tgt gag gat gcc tcg	gat gag	8352

Ser	Arg	Cys	Asn	Gln	Phe	Val	Asp	Cys	Glu	Asp	Ala	Ser	Asp	Glu		
		2625					2630					2635				
atg	aat	tgc	agt	gcc	aca	gac	tgc	agc	agc	tat	ttc	cgc	ctg	ggc	8397	
Met	Asn	Cys	Ser	Ala	Thr	Asp	Cys	Ser	Ser	Tyr	Phe	Arg	Leu	Gly		
		2640					2645					2650				
gtg	aaa	ggt	gtc	ctc	ttc	cag	ccg	tgc	gag	cgg	aca	tcc	ctg	tgc	8442	
Val	Lys	Gly	Val	Leu	Phe	Gln	Pro	Cys	Glu	Arg	Thr	Ser	Leu	Cys		
		2655					2660					2665				
tac	gca	cct	agc	tgg	gtg	tgt	gat	ggc	gcc	aac	gac	tgt	gga	gac	8487	
Tyr	Ala	Pro	Ser	Trp	Val	Cys	Asp	Gly	Ala	Asn	Asp	Cys	Gly	Asp		
		2670					2675					2680				
tac	agc	gat	gaa	cgt	gac	tgt	cca	ggc	gtg	aag	cgc	cct	agg	tgc	8532	
Tyr	Ser	Asp	Glu	Arg	Asp	Cys	Pro	Gly	Val	Lys	Arg	Pro	Arg	Cys		
		2685					2690					2695				
ccg	ctc	aat	tac	ttt	gcc	tgc	ccc	agc	ggg	cgc	tgt	atc	ccc	atg	8577	
Pro	Leu	Asn	Tyr	Phe	Ala	Cys	Pro	Ser	Gly	Arg	Cys	Ile	Pro	Met		
		2700					2705					2710				
agc	tgg	acg	tgt	gac	aag	gag	gat	gac	tgt	gag	aac	ggc	gag	gat	8622	
Ser	Trp	Thr	Cys	Asp	Lys	Glu	Asp	Asp	Cys	Glu	Asn	Gly	Glu	Asp		
		2715					2720					2725				
gag	acc	cac	tgc	aac	aag	ttc	tgc	tca	gag	gca	cag	ttc	gag	tgc	8667	
Glu	Thr	His	Cys	Asn	Lys	Phe	Cys	Ser	Glu	Ala	Gln	Phe	Glu	Cys		
		2730					2735					2740				
cag	aac	cac	cgg	tgt	atc	tcc	aag	cag	tgg	ctg	tgt	gac	ggc	agc	8712	
Gln	Asn	His	Arg	Cys	Ile	Ser	Lys	Gln	Trp	Leu	Cys	Asp	Gly	Ser		
		2745					2750					2755				
gat	gat	tgc	ggg	gat	ggc	tcc	gat	gag	gca	gct	cac	tgt	gaa	ggc	8757	
Asp	Asp	Cys	Gly	Asp	Gly	Ser	Asp	Glu	Ala	Ala	His	Cys	Glu	Gly		
		2760					2765					2770				
aag	aca	tgt	ggc	ccc	tcc	tcc	ttc	tcc	tgt	ccc	ggc	acc	cac	gtg	8802	
Lys	Thr	Cys	Gly	Pro	Ser	Ser	Phe	Ser	Cys	Pro	Gly	Thr	His	Val		
		2775					2780					2785				
tgt	gtc	cct	gag	cgc	tgg	ctc	tgt	gat	ggc	gac	aag	gac	tgt	acc	8847	
Cys	Val	Pro	Glu	Arg	Trp	Leu	Cys	Asp	Gly	Asp	Lys	Asp	Cys	Thr		
		2790					2795					2800				
gat	ggc	gcg	gat	gag	agt	gtc	act	gct	ggc	tgc	ctg	tac	aac	agc	8892	
Asp	Gly	Ala	Asp	Glu	Ser	Val	Thr	Ala	Gly	Cys	Leu	Tyr	Asn	Ser		
		2805					2810					2815				
acc	tgt	gat	gac	cgt	gag	ttc	atg	tgc	cag	aac	cgc	ttg	tgt	att	8937	
Thr	Cys	Asp	Asp	Arg	Glu	Phe	Met	Cys	Gln	Asn	Arg	Leu	Cys	Ile		
		2820					2825					2830				
ccc	aag	cat	ttc	gtg	tgc	gac	cat	gac	cgt	gac	tgt	gct	gat	ggc	8982	
Pro	Lys	His	Phe	Val	Cys	Asp	His	Asp	Arg	Asp	Cys	Ala	Asp	Gly		
		2835					2840					2845				
tct	gat	gaa	tcc	cct	gag	tgt	gag	tac	cca	acc	tgc	ggg	ccc	aat	9027	
Ser	Asp	Glu	Ser	Pro	Glu	Cys	Glu	Tyr	Pro	Thr	Cys	Gly	Pro	Asn		

2850								2855								2860								
gaa ttc cgc	tgt gcc aat ggg	cgt	tgt ctg agc tcc	cgt	cag tgg	9072																		
Glu Phe Arg	Cys Ala Asn Gly	Arg	Cys Leu Ser Ser	Arg	Gln Trp																			
2865		2870		2875																				
gaa tgt gat	ggg gag aat gac	tgt	cac gac cac agc	gat	gag gct	9117																		
Glu Cys Asp	Gly Glu Asn Asp	Cys	His Asp His Ser	Asp	Glu Ala																			
2880		2885		2890																				
ccc aag aac	cca cac tgc acc	agc	cca gag cac aaa	tgc	aat gcc	9162																		
Pro Lys Asn	Pro His Cys Thr	Ser	Pro Glu His Lys	Cys	Asn Ala																			
2895		2900		2905																				
tca tca cag	ttc ctg tgc agc	agc	ggg cgc tgc gtg	gct	gag gcg	9207																		
Ser Ser Gln	Phe Leu Cys Ser	Ser	Gly Arg Cys Val	Ala	Glu Ala																			
2910		2915		2920																				
ttg ctc tgc	aac ggc cag gac	gac	tgt ggg gac ggt	tca	gac gaa	9252																		
Leu Leu Cys	Asn Gly Gln Asp	Asp	Cys Gly Asp Gly	Ser	Asp Glu																			
2925		2930		2935																				
cgc ggg tgc	cat gtc aac gag	tgt	ctc agc cgc aag	ctc	agt ggc	9297																		
Arg Gly Cys	His Val Asn Glu	Cys	Leu Ser Arg Lys	Leu	Ser Gly																			
2940		2945		2950																				
tgc agt cag	gac tgc gag gac	ctc	aag ata ggc ttt	aag	tgc cgc	9342																		
Cys Ser Gln	Asp Cys Glu Asp	Leu	Lys Ile Gly Phe	Lys	Cys Arg																			
2955		2960		2965																				
tgt cgc ccg	ggc ttc cgg cta	aag	gac gat ggc agg	acc	tgt gcc	9387																		
Cys Arg Pro	Gly Phe Arg Leu	Lys	Asp Asp Gly Arg	Thr	Cys Ala																			
2970		2975		2980																				
gac ctg gat	gag tgc agc acc	acc	ttc ccc tgc agc	cag	ctc tgc	9432																		
Asp Leu Asp	Glu Cys Ser Thr	Thr	Phe Pro Cys Ser	Gln	Leu Cys																			
2985		2990		2995																				
atc aac acc	cac gga agt tac	aag	tgt ctg tgt gtg	gag	ggc tat	9477																		
Ile Asn Thr	His Gly Ser Tyr	Lys	Cys Leu Cys Val	Glu	Gly Tyr																			
3000		3005		3010																				
gca ccc cgt	ggc ggt gac ccc	cac	agc tgc aaa gct	gtg	acc gat	9522																		
Ala Pro Arg	Gly Gly Asp Pro	His	Ser Cys Lys Ala	Val	Thr Asp																			
3015		3020		3025																				
gag gag cca	ttt ctc atc ttt	gcc	aac cgg tac tac	ctg	cgg aag	9567																		
Glu Glu Pro	Phe Leu Ile Phe	Ala	Asn Arg Tyr Tyr	Leu	Arg Lys																			
3030		3035		3040																				
ctc aac ctg	gac ggc tcc aac	tac	aca ctg ctt aag	cag	ggc ctg	9612																		
Leu Asn Leu	Asp Gly Ser Asn	Tyr	Thr Leu Leu Lys	Gln	Gly Leu																			
3045		3050		3055																				
aac aat gcg	gtc gcc ttg gca	ttt	gac tac cga gag	cag	atg atc	9657																		
Asn Asn Ala	Val Ala Leu Ala	Phe	Asp Tyr Arg Glu	Gln	Met Ile																			
3060		3065		3070																				
tac tgg acg	ggc gtg acc acc	cag	ggc agc atg att	cgc	agg atg	9702																		
Tyr Trp Thr	Gly Val Thr Thr	Gln	Gly Ser Met Ile	Arg	Arg Met																			
3075		3080		3085																				

cac ctc aac ggc agc aac gtg cag gtt ctg cac cgg acg ggc ctt	9747
His Leu Asn Gly Ser Asn Val Gln Val Leu His Arg Thr Gly Leu	
3090 3095 3100	
agt aac cca gat ggg ctc gct gtg gac tgg gtg ggt ggc aac ctg	9792
Ser Asn Pro Asp Gly Leu Ala Val Asp Trp Val Gly Gly Asn Leu	
3105 3110 3115	
tac tgg tgt gac aag ggc aga gat acc att gag gtg tcc aag ctt	9837
Tyr Trp Cys Asp Lys Gly Arg Asp Thr Ile Glu Val Ser Lys Leu	
3120 3125 3130	
aac ggg gcc tat cgg aca gtg ctg gtc agc tct ggc ctc cgg gag	9882
Asn Gly Ala Tyr Arg Thr Val Leu Val Ser Ser Gly Leu Arg Glu	
3135 3140 3145	
ccc aga gct ctg gta gtg gat gta cag aat ggg tac ctg tac tgg	9927
Pro Arg Ala Leu Val Val Asp Val Gln Asn Gly Tyr Leu Tyr Trp	
3150 3155 3160	
aca gac tgg ggt gac cac tca ctg atc ggc cgg att ggc atg gat	9972
Thr Asp Trp Gly Asp His Ser Leu Ile Gly Arg Ile Gly Met Asp	
3165 3170 3175	
gga tct ggc cgc agc atc atc gtg gac act aag atc aca tgg ccc	10017
Gly Ser Gly Arg Ser Ile Ile Val Asp Thr Lys Ile Thr Trp Pro	
3180 3185 3190	
aat ggc ctg acc gtg gac tac gtc acg gaa cgc atc tac tgg gct	10062
Asn Gly Leu Thr Val Asp Tyr Val Thr Glu Arg Ile Tyr Trp Ala	
3195 3200 3205	
gac gcc cgt gag gac tac atc gag ttc gcc agc ctg gat ggc tcc	10107
Asp Ala Arg Glu Asp Tyr Ile Glu Phe Ala Ser Leu Asp Gly Ser	
3210 3215 3220	
aac cgt cac gtt gtg ctg agc caa gac atc cca cac atc ttt gcg	10152
Asn Arg His Val Val Leu Ser Gln Asp Ile Pro His Ile Phe Ala	
3225 3230 3235	
ctg acc cta ttt gaa gac tac gtc tac tgg aca gac tgg gaa acg	10197
Leu Thr Leu Phe Glu Asp Tyr Val Tyr Trp Thr Asp Trp Glu Thr	
3240 3245 3250	
aag tcc atc aac cgg gcc cac aag acc acg ggt gcc aac aaa aca	10242
Lys Ser Ile Asn Arg Ala His Lys Thr Thr Gly Ala Asn Lys Thr	
3255 3260 3265	
ctc ctc atc agc acc ctg cac cgg ccc atg gac tta cat gta ttc	10287
Leu Leu Ile Ser Thr Leu His Arg Pro Met Asp Leu His Val Phe	
3270 3275 3280	
cac gcc ctg cgc cag cca gat gtg ccc aat cac ccc tgc aaa gtc	10332
His Ala Leu Arg Gln Pro Asp Val Pro Asn His Pro Cys Lys Val	
3285 3290 3295	
aac aat ggt ggc tgc agc aac ctg tgc ctg ctg tcc cct ggg ggt	10377
Asn Asn Gly Gly Cys Ser Asn Leu Cys Leu Leu Ser Pro Gly Gly	
3300 3305 3310	

ggt cac aag tgc gcc tgc ccc acc aac ttc tat ctg ggt ggc gat	10422
Gly His Lys Cys Ala Cys Pro Thr Asn Phe Tyr Leu Gly Gly Asp	
3315 3320 3325	
ggc cgt acc tgt gtg tcc aac tgc aca gca agc cag ttt gtg tgc	10467
Gly Arg Thr Cys Val Ser Asn Cys Thr Ala Ser Gln Phe Val Cys	
3330 3335 3340	
aaa aat gac aag tgc atc ccc ttc tgg tgg aag tgt gac acg gag	10512
Lys Asn Asp Lys Cys Ile Pro Phe Trp Trp Lys Cys Asp Thr Glu	
3345 3350 3355	
gac gac tgt ggg gat cac tca gac gag cct cca gac tgt ccc gag	10557
Asp Asp Cys Gly Asp His Ser Asp Glu Pro Pro Asp Cys Pro Glu	
3360 3365 3370	
ttc aag tgc cgc cca ggc cag ttc cag tgc tcc acc ggc atc tgc	10602
Phe Lys Cys Arg Pro Gly Gln Phe Gln Cys Ser Thr Gly Ile Cys	
3375 3380 3385	
acc aac cct gcc ttc atc tgt gat ggg gac aat gac tgc caa gac	10647
Thr Asn Pro Ala Phe Ile Cys Asp Gly Asp Asn Asp Cys Gln Asp	
3390 3395 3400	
aat agt gac gag gcc aat tgc gac att cac gtc tgc ttg ccc agc	10692
Asn Ser Asp Glu Ala Asn Cys Asp Ile His Val Cys Leu Pro Ser	
3405 3410 3415	
caa ttc aag tgc acc aac acc aac cgc tgc att cct ggc atc ttc	10737
Gln Phe Lys Cys Thr Asn Thr Asn Arg Cys Ile Pro Gly Ile Phe	
3420 3425 3430	
cgt tgc aat ggg cag gac aac tgc ggg gac ggc gag gat gag cgg	10782
Arg Cys Asn Gly Gln Asp Asn Cys Gly Asp Gly Glu Asp Glu Arg	
3435 3440 3445	
gat tgc cct gag gtg acc tgc gcc ccc aac cag ttc cag tgc tcc	10827
Asp Cys Pro Glu Val Thr Cys Ala Pro Asn Gln Phe Gln Cys Ser	
3450 3455 3460	
atc acc aag cgc tgc atc cct cgc gtc tgg gtc tgt gac agg gat	10872
Ile Thr Lys Arg Cys Ile Pro Arg Val Trp Val Cys Asp Arg Asp	
3465 3470 3475	
aat cac tgt gtg gac ggc agt gat gag cct gcc aac tgt acc caa	10917
Asn His Cys Val Asp Gly Ser Asp Glu Pro Ala Asn Cys Thr Gln	
3480 3485 3490	
atg acc tgt gga gtg gat gag ttc cgc tgc aag gat tct ggc cgc	10962
Met Thr Cys Gly Val Asp Glu Phe Arg Cys Lys Asp Ser Gly Arg	
3495 3500 3505	
tgc atc ccc gcg cgc tgg aag tgt gac gga gaa gat gac tgt ggg	11007
Cys Ile Pro Ala Arg Trp Lys Cys Asp Gly Glu Asp Asp Cys Gly	
3510 3515 3520	
gat ggt tca gat gag ccc aag gaa gag tgt gat gag cgc acc tgt	11052
Asp Gly Ser Asp Glu Pro Lys Glu Glu Cys Asp Glu Arg Thr Cys	
3525 3530 3535	
gag cca tac cag ttc cgc tgc aaa aac aac cgc tgt gtc cca ggc	11097



Glu	Pro	Tyr	Gln	Phe	Arg	Cys	Lys	Asn	Asn	Arg	Cys	Val	Pro	Gly		
		3540					3545					3550				
cgt	tgg	caa	tgt	gac	tac	gac	aac	gac	tgc	gga	gat	aac	tcg	gac		11142
Arg	Trp	Gln	Cys	Asp	Tyr	Asp	Asn	Asp	Cys	Gly	Asp	Asn	Ser	Asp		
		3555					3560					3565				
gag	gag	agc	tgc	aca	cct	cgg	ccc	tgc	tct	gag	agt	gag	ttt	ttc		11187
Glu	Glu	Ser	Cys	Thr	Pro	Arg	Pro	Cys	Ser	Glu	Ser	Glu	Phe	Phe		
		3570					3575					3580				
tgt	gcc	aat	ggc	cgc	tgc	atc	gct	ggg	cgc	tgg	aag	tgt	gat	ggg		11232
Cys	Ala	Asn	Gly	Arg	Cys	Ile	Ala	Gly	Arg	Trp	Lys	Cys	Asp	Gly		
		3585					3590					3595				
gac	cat	gac	tgt	gcc	gac	ggc	tca	gac	gag	aaa	gac	tgc	acc	ccc		11277
Asp	His	Asp	Cys	Ala	Asp	Gly	Ser	Asp	Glu	Lys	Asp	Cys	Thr	Pro		
		3600					3605					3610				
cgc	tgt	gat	atg	gac	cag	ttc	cag	tgc	aag	agt	ggc	cac	tgc	atc		11322
Arg	Cys	Asp	Met	Asp	Gln	Phe	Gln	Cys	Lys	Ser	Gly	His	Cys	Ile		
		3615					3620					3625				
ccc	ctg	cgc	tgg	ccg	tgt	gac	gcg	gat	gct	gac	tgt	atg	gac	ggc		11367
Pro	Leu	Arg	Trp	Pro	Cys	Asp	Ala	Asp	Ala	Asp	Cys	Met	Asp	Gly		
		3630					3635					3640				
agt	gac	gag	gaa	gcc	tgt	ggc	act	ggg	gtg	agg	acc	tgc	cca	ttg		11412
Ser	Asp	Glu	Glu	Ala	Cys	Gly	Thr	Gly	Val	Arg	Thr	Cys	Pro	Leu		
		3645					3650					3655				
gat	gag	ttt	caa	tgt	aac	aac	acc	ttg	tgc	aag	ccg	ctg	gcc	tgg		11457
Asp	Glu	Phe	Gln	Cys	Asn	Asn	Thr	Leu	Cys	Lys	Pro	Leu	Ala	Trp		
		3660					3665					3670				
aag	tgt	gat	gga	gag	gac	gac	tgt	ggg	gac	aac	tca	gat	gag	aac		11502
Lys	Cys	Asp	Gly	Glu	Asp	Asp	Cys	Gly	Asp	Asn	Ser	Asp	Glu	Asn		
		3675					3680					3685				
ccc	gag	gaa	tgc	gcc	cgg	ttc	atc	tgc	cct	ccc	aac	cgg	cct	ttc		11547
Pro	Glu	Glu	Cys	Ala	Arg	Phe	Ile	Cys	Pro	Pro	Asn	Arg	Pro	Phe		
		3690					3695					3700				
cgc	tgc	aag	aat	gac	cga	gtc	tgc	ctg	tgg	att	ggg	cgc	cag	tgt		11592
Arg	Cys	Lys	Asn	Asp	Arg	Val	Cys	Leu	Trp	Ile	Gly	Arg	Gln	Cys		
		3705					3710					3715				
gat	ggc	gtg	gac	aac	tgt	gga	gat	ggg	act	gac	gag	gag	gac	tgt		11637
Asp	Gly	Val	Asp	Asn	Cys	Gly	Asp	Gly	Thr	Asp	Glu	Glu	Asp	Cys		
		3720					3725					3730				
gag	ccc	ccc	acg	gcc	cag	aac	ccc	cac	tgc	aaa	gac	aag	aag	gag		11682
Glu	Pro	Pro	Thr	Ala	Gln	Asn	Pro	His	Cys	Lys	Asp	Lys	Lys	Glu		
		3735					3740					3745				
ttc	ctg	tgc	cga	aac	cag	cgc	tgt	cta	tca	tcc	tcc	ctg	cgc	tgt		11727
Phe	Leu	Cys	Arg	Asn	Gln	Arg	Cys	Leu	Ser	Ser	Ser	Leu	Arg	Cys		
		3750					3755					3760				
aac	atg	ttc	gat	gac	tgc	ggc	gat	ggc	tcc	gat	gaa	gaa	gat	tgc		11772
Asn	Met	Phe	Asp	Asp	Cys	Gly	Asp	Gly	Ser	Asp	Glu	Glu	Asp	Cys		

3765				3770				3775				
agc atc gac	ccc aag ctg acc	agc tgt gcc acc aat	gcc agc atg	11817								
Ser Ile Asp	Pro Lys Leu Thr	Ser Cys Ala Thr Asn	Ser Met									
3780		3785	3790									
tgt ggg gac	gaa gct cgt tgt gtg	cgc act gag aaa gct	gcc tac	11862								
Cys Gly Asp	Glu Ala Arg Cys Val	Arg Thr Glu Lys Ala	Ala Tyr									
3795		3800	3805									
tgt gcc tgc	cgc tcg ggc ttc cat	act gtg ccg ggc cag	ccc gga	11907								
Cys Ala Cys	Arg Ser Gly Phe His	Thr Val Pro Gly Gln	Pro Gly									
3810		3815	3820									
tgc cag gac	atc aac gag tgc ctg	cgc ttt ggt acc tgc	tct cag	11952								
Cys Gln Asp	Ile Asn Glu Cys Leu	Arg Phe Gly Thr Cys	Ser Gln									
3825		3830	3835									
ctc tgg aac	aaa ccc aag gga ggc	cac ctc tgc agc tgt	gcc cgc	11997								
Leu Trp Asn	Lys Pro Lys Gly Gly	His Leu Cys Ser Cys	Ala Arg									
3840		3845	3850									
aac ttc atg	aag aca cac aac acc	tgc aaa gct gaa ggc	tcc gag	12042								
Asn Phe Met	Lys Thr His Asn Thr	Cys Lys Ala Glu Gly	Ser Glu									
3855		3860	3865									
tac cag gtg	cta tac atc gcg gat	gac aac gag atc cgc	agc ttg	12087								
Tyr Gln Val	Leu Tyr Ile Ala Asp	Asp Asn Glu Ile Arg	Ser Leu									
3870		3875	3880									
ttc ccg ggc	cac ccc cac tca gcc	tac gag cag aca ttc	cag ggc	12132								
Phe Pro Gly	His Pro His Ser Ala	Tyr Glu Gln Thr Phe	Gln Gly									
3885		3890	3895									
gat gag agt	gtc cgc ata gat gcc	atg gat gtc cat gtc	aag gcc	12177								
Asp Glu Ser	Val Arg Ile Asp Ala	Met Asp Val His Val	Lys Ala									
3900		3905	3910									
ggc cgt gtc	tac tgg act aac tgg	cac acg ggc aca atc	tcc tac	12222								
Gly Arg Val	Tyr Trp Thr Asn Trp	His Thr Gly Thr Ile	Ser Tyr									
3915		3920	3925									
agg agc ctg	ccc cct gcc gcc cct	cct acc act tcc aac	cgc cac	12267								
Arg Ser Leu	Pro Pro Ala Ala Pro	Pro Thr Thr Ser Asn	Arg His									
3930		3935	3940									
cgg agg cag	atc gac cgg ggt gtc	acc cac ctc aat att	tca ggg	12312								
Arg Arg Gln	Ile Asp Arg Gly Val	Thr His Leu Asn Ile	Ser Gly									
3945		3950	3955									
ctg aag atg	ccg agg ggt atc gct	atc gac tgg gtg gcc	ggg aat	12357								
Leu Lys Met	Pro Arg Gly Ile Ala	Ile Asp Trp Val Ala	Gly Asn									
3960		3965	3970									
gtg tac tgg	acc gat tcc ggc cga	gac gtg att gag gtg	gcg caa	12402								
Val Tyr Trp	Thr Asp Ser Gly Arg	Asp Val Ile Glu Val	Ala Gln									
3975		3980	3985									
atg aag ggc	gag aac cgc aag acg	ctc atc tcg ggc atg	att gat	12447								
Met Lys Gly	Glu Asn Arg Lys Thr	Leu Ile Ser Gly Met	Ile Asp									
3990		3995	4000									

gag ccc cat gcc atc gtg gtg gac cct ctg agg ggc acc atg tac	12492
Glu Pro His Ala Ile Val Val Asp Pro Leu Arg Gly Thr Met Tyr	
4005 4010 4015	
tgg tca gac tgg ggg aac cac ccc aag att gaa aca gca gcg atg	12537
Trp Ser Asp Trp Gly Asn His Pro Lys Ile Glu Thr Ala Ala Met	
4020 4025 4030	
gat ggc acc ctt cgg gag act ctc gtg caa gac aac att cag tgg	12582
Asp Gly Thr Leu Arg Glu Thr Leu Val Gln Asp Asn Ile Gln Trp	
4035 4040 4045	
cct aca ggg ctg gct gtg gac tat cac aat gaa cgg ctc tac tgg	12627
Pro Thr Gly Leu Ala Val Asp Tyr His Asn Glu Arg Leu Tyr Trp	
4050 4055 4060	
gca gat gcc aag ctt tcg gtc atc ggc agc atc cgg ctc aac ggc	12672
Ala Asp Ala Lys Leu Ser Val Ile Gly Ser Ile Arg Leu Asn Gly	
4065 4070 4075	
act gac ccc att gtg gct gct gac agc aaa cga ggc cta agt cac	12717
Thr Asp Pro Ile Val Ala Ala Asp Ser Lys Arg Gly Leu Ser His	
4080 4085 4090	
ccc ttc agc atc gat gtg ttt gaa gac tac atc tac gga gtc act	12762
Pro Phe Ser Ile Asp Val Phe Glu Asp Tyr Ile Tyr Gly Val Thr	
4095 4100 4105	
tac atc aat aat cgt gtc ttc aag atc cac aag ttt gga cac agc	12807
Tyr Ile Asn Asn Arg Val Phe Lys Ile His Lys Phe Gly His Ser	
4110 4115 4120	
ccc ttg tac aac cta act ggg ggc ctg agc cat gcc tct gat gta	12852
Pro Leu Tyr Asn Leu Thr Gly Gly Leu Ser His Ala Ser Asp Val	
4125 4130 4135	
gtc ctt tac cat caa cac aag cag cct gaa gtg acc aac ccc tgt	12897
Val Leu Tyr His Gln His Lys Gln Pro Glu Val Thr Asn Pro Cys	
4140 4145 4150	
gac cgc aag aaa tgc gaa tgg ctg tgt ctg ctg agc ccc agc ggc	12942
Asp Arg Lys Lys Cys Glu Trp Leu Cys Leu Leu Ser Pro Ser Gly	
4155 4160 4165	
cct gtc tgc acc tgt ccc aat gga aag agg ctg gat aat ggc acc	12987
Pro Val Cys Thr Cys Pro Asn Gly Lys Arg Leu Asp Asn Gly Thr	
4170 4175 4180	
tgt gtg cct gtg ccc tct cca aca ccc cct cca gat gcc cct agg	13032
Cys Val Pro Val Pro Ser Pro Thr Pro Pro Pro Asp Ala Pro Arg	
4185 4190 4195	
cct gga acc tgc act ctg cag tgc ttc aat ggt ggt agt tgt ttc	13077
Pro Gly Thr Cys Thr Leu Gln Cys Phe Asn Gly Gly Ser Cys Phe	
4200 4205 4210	
ctc aac gct cgg agg cag ccc aag tgc cgt tgc cag ccc cgt tac	13122
Leu Asn Ala Arg Arg Gln Pro Lys Cys Arg Cys Gln Pro Arg Tyr	
4215 4220 4225	

aca ggc gat	aag tgt gag ctg gat	cag tgc tgg gaa tac	tgt cac	13167
Thr Gly Asp	Lys Cys Glu Leu Asp	Gln Cys Trp Glu Tyr	Cys His	
4230	4235	4240		
aac gga ggc	acc tgt gcg gct tcc	cca tct ggc atg ccc	acg tgc	13212
Asn Gly Gly	Thr Cys Ala Ala Ser	Pro Ser Gly Met Pro	Thr Cys	
4245	4250	4255		
cgc tgt ccc	act ggc ttc acg ggc	ccc aaa tgc acc gca	cag gtg	13257
Arg Cys Pro	Thr Gly Phe Thr Gly	Pro Lys Cys Thr Ala	Gln Val	
4260	4265	4270		
tgt gca ggc	tac tgc tct aac aac	agc acc tgc acc gtc	aac cag	13302
Cys Ala Gly	Tyr Cys Ser Asn Asn	Ser Thr Cys Thr Val	Asn Gln	
4275	4280	4285		
ggc aac cag	ccc cag tgc cga tgt	cta cct ggc ttc ctg	ggc gac	13347
Gly Asn Gln	Pro Gln Cys Arg Cys	Leu Pro Gly Phe Leu	Gly Asp	
4290	4295	4300		
cgt tgc cag	tac cgg cag tgc tct	ggc ttc tgt gag aac	ttt ggc	13392
Arg Cys Gln	Tyr Arg Gln Cys Ser	Gly Phe Cys Glu Asn	Phe Gly	
4305	4310	4315		
acc tgt cag	atg gct gct gat ggc	tcc cga caa tgt cgc	tgc acc	13437
Thr Cys Gln	Met Ala Ala Asp Gly	Ser Arg Gln Cys Arg	Cys Thr	
4320	4325	4330		
gtc tac ttt	gag gga cca agg tgt	gag gtg aac aag tgt	agt cgc	13482
Val Tyr Phe	Glu Gly Pro Arg Cys	Glu Val Asn Lys Cys	Ser Arg	
4335	4340	4345		
tgt ctc caa	ggc gcc tgt gtg gtc	aat aag cag acc gga	gat gtc	13527
Cys Leu Gln	Gly Ala Cys Val Val	Asn Lys Gln Thr Gly	Asp Val	
4350	4355	4360		
aca tgc aac	tgc act gat ggc cgg	gta gcc ccc agt tgt	ctc acc	13572
Thr Cys Asn	Cys Thr Asp Gly Arg	Val Ala Pro Ser Cys	Leu Thr	
4365	4370	4375		
tgc atc gat	cac tgt agc aat ggt	ggc tcc tgc acc atg	aac agc	13617
Cys Ile Asp	His Cys Ser Asn Gly	Gly Ser Cys Thr Met	Asn Ser	
4380	4385	4390		
aag atg atg	cct gag tgc cag tgc	ccg ccc cat atg aca	gga ccc	13662
Lys Met Met	Pro Glu Cys Gln Cys	Pro Pro His Met Thr	Gly Pro	
4395	4400	4405		
cgg tgc cag	gag cag gtt gtt agt	cag caa cag cct ggg	cat atg	13707
Arg Cys Gln	Glu Gln Val Val Ser	Gln Gln Gln Pro Gly	His Met	
4410	4415	4420		
gcc tcc atc	ctg atc cct ctg ctg	ctg ctt ctc ctg ctg	ctt ctg	13752
Ala Ser Ile	Leu Ile Pro Leu Leu	Leu Leu Leu Leu Leu	Leu Leu	
4425	4430	4435		
gtg gct ggc	gtg gtg ttc tgg tat	aag cgg cga gtc cga	ggg gct	13797
Val Ala Gly	Val Val Phe Trp Tyr	Lys Arg Arg Val Arg	Gly Ala	
4440	4445	4450		
aag ggc ttc	cag cac cag cgg atg	acc aat ggg gcc atg	aat gtg	13842

Lys Gly Phe	Gln His Gln Arg Met	Thr Asn Gly Ala Met	Asn Val	
4455	4460	4465		
gaa att gga	aac cct acc tac aag	atg tat gaa ggt gga	gag ccc	13887
Glu Ile Gly	Asn Pro Thr Tyr Lys	Met Tyr Glu Gly Gly	Glu Pro	
4470	4475	4480		
gat gat gtc	ggg ggc cta ctg gat	gct gat ttt gcc ctt	gac cct	13932
Asp Asp Val	Gly Gly Leu Leu Asp	Ala Asp Phe Ala Leu	Asp Pro	
4485	4490	4495		
gac aag cct	acc aac ttc acc aac	cca gtg tat gcc acg	ctc tac	13977
Asp Lys Pro	Thr Asn Phe Thr Asn	Pro Val Tyr Ala Thr	Leu Tyr	
4500	4505	4510		
atg ggg ggc	cac ggc agc cgc cat	tcc ctg gcc agc acg	gac gag	14022
Met Gly Gly	His Gly Ser Arg His	Ser Leu Ala Ser Thr	Asp Glu	
4515	4520	4525		
aag cga gaa	ctg ctg ggc cgg gga	cct gaa gac gag ata	gga gat	14067
Lys Arg Glu	Leu Leu Gly Arg Gly	Pro Glu Asp Glu Ile	Gly Asp	
4530	4535	4540		
ccc ttg gca	tag ggccctgccc cgacggatgt	ccccagaaag cccctgcca		14119
Pro Leu Ala				
4545				
catgagtctt	tcaatgaacc ccctccccag	cgggcccttc tccggccctg	ccgggtgtac	14179
aaatgtaaaa	atgaaggaat tactttttat	atgtgagcga gcaagcgagc	aagcacagta	14239
ttatctcttt	gcatttcctt cctgcctgct	cctcagtatc ccccccatgc	tgcccttgagg	14299
ggggcggggag	ggcttttgtg ctcaaaggta	tgaaggagtc cacatgttcc	ctaccgagca	14359
tacccttgga	agcctggcgg cacggcctcc	ccaccacgcc tgtgcaagac	actcaacggg	14419
gctccgtgtc	ccagctttcc tttccttggc	tctctggggg tagttcaggg	gaggtggagt	14479
cctctgctga	ccctgtctgg aagatttggc	tctagctgag gaaggagtct	tttagttgag	14539
ggaagtcacc	ccaaacccca gctcccactt	tcaggggcac ctctcagatg	gccatgctca	14599
gtatcccttc	cagacaggcc ctcccctctc	tagcgcccc tctgtggctc	ctagggctga	14659
acacattctt	tggtaaactgt cccccaagcc	tcccatcccc ctgagggcca	ggaagagtcg	14719
gggcacacca	aggaagggca agcgggcagc	cccattttgg ggacgtgaac	gttttaataa	14779
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<211>	4545			
<212>	PRT			
<213>	Mus sp.			
<400>	2			

Met	Leu	Thr	Pro	Pro	Leu	Leu	Leu	Leu	Val	Pro	Leu	Leu	Ser	Ala	Leu	1	5	10	15
Val	Ser	Gly	Ala	Thr	Met	Asp	Ala	Pro	Lys	Thr	Cys	Ser	Pro	Lys	Gln	20	25	30	
Phe	Ala	Cys	Arg	Asp	Gln	Ile	Thr	Cys	Ile	Ser	Lys	Gly	Trp	Arg	Cys	35	40	45	
Asp	Gly	Glu	Arg	Asp	Cys	Pro	Asp	Gly	Ser	Asp	Glu	Ala	Pro	Glu	Ile	50	55	60	
Cys	Pro	Gln	Ser	Lys	Ala	Gln	Arg	Cys	Pro	Pro	Asn	Glu	His	Ser	Cys	65	70	75	80
Leu	Gly	Thr	Glu	Leu	Cys	Val	Pro	Met	Ser	Arg	Leu	Cys	Asn	Gly	Ile	85	90	95	
Gln	Asp	Cys	Met	Asp	Gly	Ser	Asp	Glu	Gly	Ala	His	Cys	Arg	Glu	Leu	100	105	110	
Arg	Ala	Asn	Cys	Ser	Arg	Met	Gly	Cys	Gln	His	His	Cys	Val	Pro	Thr	115	120	125	
Pro	Ser	Gly	Pro	Thr	Cys	Tyr	Cys	Asn	Ser	Ser	Phe	Gln	Leu	Glu	Ala	130	135	140	
Asp	Gly	Lys	Thr	Cys	Lys	Asp	Phe	Asp	Glu	Cys	Ser	Val	Tyr	Gly	Thr	145	150	155	160
Cys	Ser	Gln	Leu	Cys	Thr	Asn	Thr	Asp	Gly	Ser	Phe	Thr	Cys	Gly	Cys	165	170	175	
Val	Glu	Gly	Tyr	Leu	Leu	Gln	Pro	Asp	Asn	Arg	Ser	Cys	Lys	Ala	Lys	180	185	190	
Asn	Glu	Pro	Val	Asp	Arg	Pro	Pro	Val	Leu	Leu	Ile	Ala	Asn	Ser	Gln	195	200	205	
Asn	Ile	Leu	Ala	Thr	Tyr	Leu	Ser	Gly	Ala	Gln	Val	Ser	Thr	Ile	Thr	210	215	220	
Pro	Thr	Ser	Thr	Arg	Gln	Thr	Thr	Ala	Met	Asp	Phe	Ser	Tyr	Ala	Asn	225	230	235	240
Glu	Thr	Val	Cys	Trp	Val	His	Val	Gly	Asp	Ser	Ala	Ala	Gln	Thr	Gln				

245								250				255			
Leu	Lys	Cys	Ala	Arg	Met	Pro	Gly	Leu	Lys	Gly	Phe	Val	Asp	Glu	His
			260					265					270		
Thr	Ile	Asn	Ile	Ser	Leu	Ser	Leu	His	His	Val	Glu	Gln	Met	Ala	Ile
		275					280					285			
Asp	Trp	Leu	Thr	Gly	Asn	Phe	Tyr	Phe	Val	Asp	Asp	Ile	Asp	Asp	Arg
	290					295					300				
Ile	Phe	Val	Cys	Asn	Arg	Asn	Gly	Asp	Thr	Cys	Val	Thr	Leu	Leu	Asp
305					310					315					320
Leu	Glu	Leu	Tyr	Asn	Pro	Lys	Gly	Ile	Ala	Leu	Asp	Pro	Ala	Met	Gly
				325					330					335	
Lys	Val	Phe	Phe	Thr	Asp	Tyr	Gly	Gln	Ile	Pro	Lys	Val	Glu	Arg	Cys
			340					345					350		
Asp	Met	Asp	Gly	Gln	Asn	Arg	Thr	Lys	Leu	Val	Asp	Ser	Lys	Ile	Val
		355					360					365			
Phe	Pro	His	Gly	Ile	Thr	Leu	Asp	Leu	Val	Ser	Arg	Leu	Val	Tyr	Trp
	370					375					380				
Ala	Asp	Ala	Tyr	Leu	Asp	Tyr	Ile	Glu	Val	Val	Asp	Tyr	Glu	Gly	Lys
385					390					395					400
Gly	Arg	Gln	Thr	Ile	Ile	Gln	Gly	Ile	Leu	Ile	Glu	His	Leu	Tyr	Gly
				405					410					415	
Leu	Thr	Val	Phe	Glu	Asn	Tyr	Leu	Tyr	Ala	Thr	Asn	Ser	Asp	Asn	Ala
			420					425					430		
Asn	Thr	Gln	Gln	Lys	Thr	Ser	Val	Ile	Arg	Val	Asn	Arg	Phe	Asn	Ser
		435					440					445			
Thr	Glu	Tyr	Gln	Val	Val	Thr	Arg	Val	Asp	Lys	Gly	Gly	Ala	Leu	His
	450					455					460				
Ile	Tyr	His	Gln	Arg	Arg	Gln	Pro	Arg	Val	Arg	Ser	His	Ala	Cys	Glu
465					470					475					480
Asn	Asp	Gln	Tyr	Gly	Lys	Pro	Gly	Gly	Cys	Ser	Asp	Ile	Cys	Leu	Leu
				485					490					495	

Ala	Asn	Ser	His	Lys	Ala	Arg	Thr	Cys	Arg	Cys	Arg	Ser	Gly	Phe	Ser	500	505	510
Leu	Gly	Ser	Asp	Gly	Lys	Ser	Cys	Lys	Lys	Pro	Glu	His	Glu	Leu	Phe	515	520	525
Leu	Val	Tyr	Gly	Lys	Gly	Arg	Pro	Gly	Ile	Ile	Arg	Gly	Met	Asp	Met	530	535	540
Gly	Ala	Lys	Val	Pro	Asp	Glu	His	Met	Ile	Pro	Ile	Glu	Asn	Leu	Met	545	550	555
Asn	Pro	Arg	Ala	Leu	Asp	Phe	His	Ala	Glu	Thr	Gly	Phe	Ile	Tyr	Phe	565	570	575
Ala	Asp	Thr	Thr	Ser	Tyr	Leu	Ile	Gly	Arg	Gln	Lys	Ile	Asp	Gly	Thr	580	585	590
Glu	Arg	Glu	Thr	Ile	Leu	Lys	Asp	Gly	Ile	His	Asn	Val	Glu	Gly	Val	595	600	605
Ala	Val	Asp	Trp	Met	Gly	Asp	Asn	Leu	Tyr	Trp	Thr	Asp	Asp	Gly	Pro	610	615	620
Lys	Lys	Thr	Ile	Ser	Val	Ala	Arg	Leu	Glu	Lys	Ala	Ala	Gln	Thr	Arg	625	630	635
Lys	Thr	Leu	Ile	Glu	Gly	Lys	Met	Thr	His	Pro	Arg	Ala	Ile	Val	Val	645	650	655
Asp	Pro	Leu	Asn	Gly	Trp	Met	Tyr	Trp	Thr	Asp	Trp	Glu	Glu	Asp	Pro	660	665	670
Lys	Asp	Ser	Arg	Arg	Gly	Arg	Leu	Glu	Arg	Ala	Trp	Met	Asp	Gly	Ser	675	680	685
His	Arg	Asp	Ile	Phe	Val	Thr	Ser	Lys	Thr	Val	Leu	Trp	Pro	Asn	Gly	690	695	700
Leu	Ser	Leu	Asp	Ile	Pro	Ala	Gly	Arg	Leu	Tyr	Trp	Val	Asp	Ala	Phe	705	710	715
Tyr	Asp	Arg	Ile	Glu	Thr	Ile	Leu	Leu	Asn	Gly	Thr	Asp	Arg	Lys	Ile	725	730	735



Val Tyr Glu Gly Pro Glu Leu Asn His Ala Phe Gly Leu Cys His His  
740 745 750

Gly Asn Tyr Leu Phe Trp Thr Glu Tyr Arg Ser Gly Ser Val Tyr Arg  
755 760 765

Leu Glu Arg Gly Val Ala Gly Ala Pro Pro Thr Val Thr Leu Leu Arg  
770 775 780

Ser Glu Arg Pro Pro Ile Phe Glu Ile Arg Met Tyr Asp Ala His Glu  
785 790 795 800

Gln Gln Val Gly Thr Asn Lys Cys Arg Val Asn Asn Gly Gly Cys Ser  
805 810 815

Ser Leu Cys Leu Ala Thr Pro Gly Ser Arg Gln Cys Ala Cys Ala Glu  
820 825 830

Asp Gln Val Leu Asp Thr Asp Gly Val Thr Cys Leu Ala Asn Pro Ser  
835 840 845

Tyr Val Pro Pro Pro Gln Cys Gln Pro Gly Gln Phe Ala Cys Ala Asn  
850 855 860

Asn Arg Cys Ile Gln Glu Arg Trp Lys Cys Asp Gly Asp Asn Asp Cys  
865 870 875 880

Leu Asp Asn Ser Asp Glu Ala Pro Ala Leu Cys His Gln His Thr Cys  
885 890 895

Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn Arg  
900 905 910

Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu Ser  
915 920 925

Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser Cys  
930 935 940

Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp Asp  
945 950 955 960

Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro Thr  
965 970 975

Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile Asn  
980 985 990

Ile Asn Trp Arg Cys Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser Asp  
995 1000 1005

Glu Ala Gly Cys Ser His Ser Cys Ser Ser Thr Gln Phe Lys Cys  
1010 1015 1020

Asn Ser Gly Arg Cys Ile Pro Glu His Trp Thr Cys Asp Gly Asp  
1025 1030 1035

Asn Asp Cys Gly Asp Tyr Ser Asp Glu Thr His Ala Asn Cys Thr  
1040 1045 1050

Asn Gln Ala Thr Arg Pro Pro Gly Gly Cys His Ser Asp Glu Phe  
1055 1060 1065

Gln Cys Pro Leu Asp Gly Leu Cys Ile Pro Leu Arg Trp Arg Cys  
1070 1075 1080

Asp Gly Asp Thr Asp Cys Met Asp Ser Ser Asp Glu Lys Ser Cys  
1085 1090 1095

Glu Gly Val Thr His Val Cys Asp Pro Asn Val Lys Phe Gly Cys  
1100 1105 1110

Lys Asp Ser Ala Arg Cys Ile Ser Lys Ala Trp Val Cys Asp Gly  
1115 1120 1125

Asp Ser Asp Cys Glu Asp Asn Ser Asp Glu Glu Asn Cys Glu Ala  
1130 1135 1140

Leu Ala Cys Arg Pro Pro Ser His Pro Cys Ala Asn Asn Thr Ser  
1145 1150 1155

Val Cys Leu Pro Pro Asp Lys Leu Cys Asp Gly Lys Asp Asp Cys  
1160 1165 1170

Gly Asp Gly Ser Asp Glu Gly Glu Leu Cys Asp Gln Cys Ser Leu  
1175 1180 1185

Asn Asn Gly Gly Cys Ser His Asn Cys Ser Val Ala Pro Gly Glu  
1190 1195 1200

Gly Ile Val Cys Ser Cys Pro Leu Gly Met Glu Leu Gly Ser Asp

1205		1210		1215
Asn His Thr Cys Gln Ile Gln Ser Tyr Cys Ala Lys His Leu Lys	1220	1225	1230	
Cys Ser Gln Lys Cys Asp Gln Asn Lys Phe Ser Val Lys Cys Ser	1235	1240	1245	
Cys Tyr Glu Gly Trp Val Leu Glu Pro Asp Gly Glu Thr Cys Arg	1250	1255	1260	
Ser Leu Asp Pro Phe Lys Leu Phe Ile Ile Phe Ser Asn Arg His	1265	1270	1275	
Glu Ile Arg Arg Ile Asp Leu His Lys Gly Asp Tyr Ser Val Leu	1280	1285	1290	
Val Pro Gly Leu Arg Asn Thr Ile Ala Leu Asp Phe His Leu Ser	1295	1300	1305	
Gln Ser Ala Leu Tyr Trp Thr Asp Ala Val Glu Asp Lys Ile Tyr	1310	1315	1320	
Arg Gly Lys Leu Leu Asp Asn Gly Ala Leu Thr Ser Phe Glu Val	1325	1330	1335	
Val Ile Gln Tyr Gly Leu Ala Thr Pro Glu Gly Leu Ala Val Asp	1340	1345	1350	
Trp Ile Ala Gly Asn Ile Tyr Trp Val Glu Ser Asn Leu Asp Gln	1355	1360	1365	
Ile Glu Val Ala Lys Leu Asp Gly Thr Leu Arg Thr Thr Leu Leu	1370	1375	1380	
Ala Gly Asp Ile Glu His Pro Arg Ala Ile Ala Leu Asp Pro Arg	1385	1390	1395	
Asp Gly Ile Leu Phe Trp Thr Asp Trp Asp Ala Ser Leu Pro Arg	1400	1405	1410	
Ile Glu Ala Ala Ser Met Ser Gly Ala Gly Arg Arg Thr Ile His	1415	1420	1425	
Arg Glu Thr Gly Ser Gly Gly Cys Ala Asn Gly Leu Thr Val Asp				

1430		1435		1440
Tyr Leu Glu Lys Arg Ile Leu Trp Ile Asp Ala Arg Ser Asp Ala	1445	1450		1455
Ile Tyr Ser Ala Arg Tyr Asp Gly Ser Gly His Met Glu Val Leu	1460	1465		1470
Arg Gly His Glu Phe Leu Ser His Pro Phe Ala Val Thr Leu Tyr	1475	1480		1485
Gly Gly Glu Val Tyr Trp Thr Asp Trp Arg Thr Asn Thr Leu Ala	1490	1495		1500
Lys Ala Asn Lys Trp Thr Gly His Asn Val Thr Val Val Gln Arg	1505	1510		1515
Thr Asn Thr Gln Pro Phe Asp Leu Gln Val Tyr His Pro Ser Arg	1520	1525		1530
Gln Pro Met Ala Pro Asn Pro Cys Glu Ala Asn Gly Gly Arg Gly	1535	1540		1545
Pro Cys Ser His Leu Cys Leu Ile Asn Tyr Asn Arg Thr Val Ser	1550	1555		1560
Trp Ala Cys Pro His Leu Met Lys Leu His Lys Asp Asn Thr Thr	1565	1570		1575
Cys Tyr Glu Phe Lys Lys Phe Leu Leu Tyr Ala Arg Gln Met Glu	1580	1585		1590
Ile Arg Gly Val Asp Leu Asp Ala Pro Tyr Tyr Asn Tyr Ile Ile	1595	1600		1605
Ser Phe Thr Val Pro Asp Ile Asp Asn Val Thr Val Leu Asp Tyr	1610	1615		1620
Asp Ala Arg Glu Gln Arg Val Tyr Trp Ser Asp Val Arg Thr Gln	1625	1630		1635
Ala Ile Lys Arg Ala Phe Ile Asn Gly Thr Gly Val Glu Thr Val	1640	1645		1650
Val Ser Ala Asp Leu Pro Asn Ala His Gly Leu Ala Val Asp Trp	1655	1660		1665

Val	Ser	Arg	Asn	Leu	Phe	Trp	Thr	Ser	Tyr	Asp	Thr	Asn	Lys	Lys
1670						1675					1680			
Gln	Ile	Asn	Val	Ala	Arg	Leu	Asp	Gly	Ser	Phe	Lys	Asn	Ala	Val
1685						1690					1695			
Val	Gln	Gly	Leu	Glu	Gln	Pro	His	Gly	Leu	Val	Val	His	Pro	Leu
1700						1705					1710			
Arg	Gly	Lys	Leu	Tyr	Trp	Thr	Asp	Gly	Asp	Asn	Ile	Ser	Met	Ala
1715						1720					1725			
Asn	Met	Asp	Gly	Ser	Asn	His	Thr	Leu	Leu	Phe	Ser	Gly	Gln	Lys
1730						1735					1740			
Gly	Pro	Val	Gly	Leu	Ala	Ile	Asp	Phe	Pro	Glu	Ser	Lys	Leu	Tyr
1745						1750					1755			
Trp	Ile	Ser	Ser	Gly	Asn	His	Thr	Ile	Asn	Arg	Cys	Asn	Leu	Asp
1760						1765					1770			
Gly	Ser	Glu	Leu	Glu	Val	Ile	Asp	Thr	Met	Arg	Ser	Gln	Leu	Gly
1775						1780					1785			
Lys	Ala	Thr	Ala	Leu	Ala	Ile	Met	Gly	Asp	Lys	Leu	Trp	Trp	Ala
1790						1795					1800			
Asp	Gln	Val	Ser	Glu	Lys	Met	Gly	Thr	Cys	Asn	Lys	Ala	Asp	Gly
1805						1810					1815			
Ser	Gly	Ser	Val	Val	Leu	Arg	Asn	Ser	Thr	Thr	Leu	Val	Met	His
1820						1825					1830			
Met	Lys	Val	Tyr	Asp	Glu	Ser	Ile	Gln	Leu	Glu	His	Glu	Gly	Thr
1835						1840					1845			
Asn	Pro	Cys	Ser	Val	Asn	Asn	Gly	Asp	Cys	Ser	Gln	Leu	Cys	Leu
1850						1855					1860			
Pro	Thr	Ser	Glu	Thr	Thr	Arg	Ser	Cys	Met	Cys	Thr	Ala	Gly	Tyr
1865						1870					1875			
Ser	Leu	Arg	Ser	Gly	Gln	Gln	Ala	Cys	Glu	Gly	Val	Gly	Ser	Phe
1880						1885					1890			

Leu	Leu	Tyr	Ser	Val	His	Glu	Gly	Ile	Arg	Gly	Ile	Pro	Leu	Asp
1895						1900					1905			
Pro	Asn	Asp	Lys	Ser	Asp	Ala	Leu	Val	Pro	Val	Ser	Gly	Thr	Ser
1910						1915					1920			
Leu	Ala	Val	Gly	Ile	Asp	Phe	His	Ala	Glu	Asn	Asp	Thr	Ile	Tyr
1925						1930					1935			
Trp	Val	Asp	Met	Gly	Leu	Ser	Thr	Ile	Ser	Arg	Ala	Lys	Arg	Asp
1940						1945					1950			
Gln	Thr	Trp	Arg	Glu	Asp	Val	Val	Thr	Asn	Gly	Ile	Gly	Arg	Val
1955						1960					1965			
Glu	Gly	Ile	Ala	Val	Asp	Trp	Ile	Ala	Gly	Asn	Ile	Tyr	Trp	Thr
1970						1975					1980			
Asp	Gln	Gly	Phe	Asp	Val	Ile	Glu	Val	Ala	Arg	Leu	Asn	Gly	Ser
1985						1990					1995			
Phe	Arg	Tyr	Val	Val	Ile	Ser	Gln	Gly	Leu	Asp	Lys	Pro	Arg	Ala
2000						2005					2010			
Ile	Thr	Val	His	Pro	Glu	Lys	Gly	Tyr	Leu	Phe	Trp	Thr	Glu	Trp
2015						2020					2025			
Gly	His	Tyr	Pro	Arg	Ile	Glu	Arg	Ser	Arg	Leu	Asp	Gly	Thr	Glu
2030						2035					2040			
Arg	Val	Val	Leu	Val	Asn	Val	Ser	Ile	Ser	Trp	Pro	Asn	Gly	Ile
2045						2050					2055			
Ser	Val	Asp	Tyr	Gln	Gly	Gly	Lys	Leu	Tyr	Trp	Cys	Asp	Ala	Arg
2060						2065					2070			
Met	Asp	Lys	Ile	Glu	Arg	Ile	Asp	Leu	Glu	Thr	Gly	Glu	Asn	Arg
2075						2080					2085			
Glu	Val	Val	Leu	Ser	Ser	Asn	Asn	Met	Asp	Met	Phe	Ser	Val	Ser
2090						2095					2100			
Val	Phe	Glu	Asp	Phe	Ile	Tyr	Trp	Ser	Asp	Arg	Thr	His	Ala	Asn
2105						2110					2115			

Gly	Ser	Ile	Lys	Arg	Gly	Cys	Lys	Asp	Asn	Ala	Thr	Asp	Ser	Val
2120						2125					2130			
Pro	Leu	Arg	Thr	Gly	Ile	Gly	Val	Gln	Leu	Lys	Asp	Ile	Lys	Val
2135						2140					2145			
Phe	Asn	Arg	Asp	Arg	Gln	Lys	Gly	Thr	Asn	Val	Cys	Ala	Val	Ala
2150						2155					2160			
Asn	Gly	Gly	Cys	Gln	Gln	Leu	Cys	Leu	Tyr	Arg	Gly	Gly	Gly	Gln
2165						2170					2175			
Arg	Ala	Cys	Ala	Cys	Ala	His	Gly	Met	Leu	Ala	Glu	Asp	Gly	Ala
2180						2185					2190			
Ser	Cys	Arg	Glu	Tyr	Ala	Gly	Tyr	Leu	Leu	Tyr	Ser	Glu	Arg	Thr
2195						2200					2205			
Ile	Leu	Lys	Ser	Ile	His	Leu	Ser	Asp	Glu	Arg	Asn	Leu	Asn	Ala
2210						2215					2220			
Pro	Val	Gln	Pro	Phe	Glu	Asp	Pro	Glu	His	Met	Lys	Asn	Val	Ile
2225						2230					2235			
Ala	Leu	Ala	Phe	Asp	Tyr	Arg	Ala	Gly	Thr	Ser	Pro	Gly	Thr	Pro
2240						2245					2250			
Asn	Arg	Ile	Phe	Phe	Ser	Asp	Ile	His	Phe	Gly	Asn	Ile	Gln	Gln
2255						2260					2265			
Ile	Asn	Asp	Asp	Gly	Ser	Gly	Arg	Thr	Thr	Ile	Val	Glu	Asn	Val
2270						2275					2280			
Gly	Ser	Val	Glu	Gly	Leu	Ala	Tyr	His	Arg	Gly	Trp	Asp	Thr	Leu
2285						2290					2295			
Tyr	Trp	Thr	Ser	Tyr	Thr	Thr	Ser	Thr	Ile	Thr	Arg	His	Thr	Val
2300						2305					2310			
Asp	Gln	Thr	Arg	Pro	Gly	Ala	Phe	Glu	Arg	Glu	Thr	Val	Ile	Thr
2315						2320					2325			
Met	Ser	Gly	Asp	Asp	His	Pro	Arg	Ala	Phe	Val	Leu	Asp	Glu	Cys
2330						2335					2340			
Gln	Asn	Leu	Met	Phe	Trp	Thr	Asn	Trp	Asn	Glu	Leu	His	Pro	Ser

2345		2350		2355
Ile Met Arg Ala Ala Leu Ser Gly Ala Asn Val Leu Thr Leu Ile				
2360		2365		2370
Glu Lys Asp Ile Arg Thr Pro Asn Gly Leu Ala Ile Asp His Arg				
2375		2380		2385
Ala Glu Lys Leu Tyr Phe Ser Asp Ala Thr Leu Asp Lys Ile Glu				
2390		2395		2400
Arg Cys Glu Tyr Asp Gly Ser His Arg Tyr Val Ile Leu Lys Ser				
2405		2410		2415
Glu Pro Val His Pro Phe Gly Leu Ala Val Tyr Gly Glu His Ile				
2420		2425		2430
Phe Trp Thr Asp Trp Val Arg Arg Ala Val Gln Arg Ala Asn Lys				
2435		2440		2445
Tyr Val Gly Ser Asp Met Lys Leu Leu Arg Val Asp Ile Pro Gln				
2450		2455		2460
Gln Pro Met Gly Ile Ile Ala Val Ala Asn Asp Thr Asn Ser Cys				
2465		2470		2475
Glu Leu Ser Pro Cys Arg Ile Asn Asn Gly Gly Cys Gln Asp Leu				
2480		2485		2490
Cys Leu Leu Thr His Gln Gly His Val Asn Cys Ser Cys Arg Gly				
2495		2500		2505
Gly Arg Ile Leu Gln Glu Asp Phe Thr Cys Arg Ala Val Asn Ser				
2510		2515		2520
Ser Cys Arg Ala Gln Asp Glu Phe Glu Cys Ala Asn Gly Glu Cys				
2525		2530		2535
Ile Ser Phe Ser Leu Thr Cys Asp Gly Val Ser His Cys Lys Asp				
2540		2545		2550
Lys Ser Asp Glu Lys Pro Ser Tyr Cys Asn Ser Arg Arg Cys Lys				
2555		2560		2565
Lys Thr Phe Arg Gln Cys Asn Asn Gly Arg Cys Val Ser Asn Met				
2570		2575		2580



Leu	Trp	Cys	Asn	Gly	Val	Asp	Tyr	Cys	Gly	Asp	Gly	Ser	Asp	Glu
2585						2590					2595			
Ile	Pro	Cys	Asn	Lys	Thr	Ala	Cys	Gly	Val	Gly	Glu	Phe	Arg	Cys
2600						2605					2610			
Arg	Asp	Gly	Ser	Cys	Ile	Gly	Asn	Ser	Ser	Arg	Cys	Asn	Gln	Phe
2615						2620					2625			
Val	Asp	Cys	Glu	Asp	Ala	Ser	Asp	Glu	Met	Asn	Cys	Ser	Ala	Thr
2630						2635					2640			
Asp	Cys	Ser	Ser	Tyr	Phe	Arg	Leu	Gly	Val	Lys	Gly	Val	Leu	Phe
2645						2650					2655			
Gln	Pro	Cys	Glu	Arg	Thr	Ser	Leu	Cys	Tyr	Ala	Pro	Ser	Trp	Val
2660						2665					2670			
Cys	Asp	Gly	Ala	Asn	Asp	Cys	Gly	Asp	Tyr	Ser	Asp	Glu	Arg	Asp
2675						2680					2685			
Cys	Pro	Gly	Val	Lys	Arg	Pro	Arg	Cys	Pro	Leu	Asn	Tyr	Phe	Ala
2690						2695					2700			
Cys	Pro	Ser	Gly	Arg	Cys	Ile	Pro	Met	Ser	Trp	Thr	Cys	Asp	Lys
2705						2710					2715			
Glu	Asp	Asp	Cys	Glu	Asn	Gly	Glu	Asp	Glu	Thr	His	Cys	Asn	Lys
2720						2725					2730			
Phe	Cys	Ser	Glu	Ala	Gln	Phe	Glu	Cys	Gln	Asn	His	Arg	Cys	Ile
2735						2740					2745			
Ser	Lys	Gln	Trp	Leu	Cys	Asp	Gly	Ser	Asp	Asp	Cys	Gly	Asp	Gly
2750						2755					2760			
Ser	Asp	Glu	Ala	Ala	His	Cys	Glu	Gly	Lys	Thr	Cys	Gly	Pro	Ser
2765						2770					2775			
Ser	Phe	Ser	Cys	Pro	Gly	Thr	His	Val	Cys	Val	Pro	Glu	Arg	Trp
2780						2785					2790			
Leu	Cys	Asp	Gly	Asp	Lys	Asp	Cys	Thr	Asp	Gly	Ala	Asp	Glu	Ser
2795						2800					2805			

Val	Thr	Ala	Gly	Cys	Leu	Tyr	Asn	Ser	Thr	Cys	Asp	Asp	Arg	Glu
2810						2815					2820			
Phe	Met	Cys	Gln	Asn	Arg	Leu	Cys	Ile	Pro	Lys	His	Phe	Val	Cys
2825						2830					2835			
Asp	His	Asp	Arg	Asp	Cys	Ala	Asp	Gly	Ser	Asp	Glu	Ser	Pro	Glu
2840						2845					2850			
Cys	Glu	Tyr	Pro	Thr	Cys	Gly	Pro	Asn	Glu	Phe	Arg	Cys	Ala	Asn
2855						2860					2865			
Gly	Arg	Cys	Leu	Ser	Ser	Arg	Gln	Trp	Glu	Cys	Asp	Gly	Glu	Asn
2870						2875					2880			
Asp	Cys	His	Asp	His	Ser	Asp	Glu	Ala	Pro	Lys	Asn	Pro	His	Cys
2885						2890					2895			
Thr	Ser	Pro	Glu	His	Lys	Cys	Asn	Ala	Ser	Ser	Gln	Phe	Leu	Cys
2900						2905					2910			
Ser	Ser	Gly	Arg	Cys	Val	Ala	Glu	Ala	Leu	Leu	Cys	Asn	Gly	Gln
2915						2920					2925			
Asp	Asp	Cys	Gly	Asp	Gly	Ser	Asp	Glu	Arg	Gly	Cys	His	Val	Asn
2930						2935					2940			
Glu	Cys	Leu	Ser	Arg	Lys	Leu	Ser	Gly	Cys	Ser	Gln	Asp	Cys	Glu
2945						2950					2955			
Asp	Leu	Lys	Ile	Gly	Phe	Lys	Cys	Arg	Cys	Arg	Pro	Gly	Phe	Arg
2960						2965					2970			
Leu	Lys	Asp	Asp	Gly	Arg	Thr	Cys	Ala	Asp	Leu	Asp	Glu	Cys	Ser
2975						2980					2985			
Thr	Thr	Phe	Pro	Cys	Ser	Gln	Leu	Cys	Ile	Asn	Thr	His	Gly	Ser
2990						2995					3000			
Tyr	Lys	Cys	Leu	Cys	Val	Glu	Gly	Tyr	Ala	Pro	Arg	Gly	Gly	Asp
3005						3010					3015			
Pro	His	Ser	Cys	Lys	Ala	Val	Thr	Asp	Glu	Glu	Pro	Phe	Leu	Ile
3020						3025					3030			

Phe	Ala	Asn	Arg	Tyr	Tyr	Leu	Arg	Lys	Leu	Asn	Leu	Asp	Gly	Ser
3035						3040					3045			
Asn	Tyr	Thr	Leu	Leu	Lys	Gln	Gly	Leu	Asn	Asn	Ala	Val	Ala	Leu
3050						3055					3060			
Ala	Phe	Asp	Tyr	Arg	Glu	Gln	Met	Ile	Tyr	Trp	Thr	Gly	Val	Thr
3065						3070					3075			
Thr	Gln	Gly	Ser	Met	Ile	Arg	Arg	Met	His	Leu	Asn	Gly	Ser	Asn
3080						3085					3090			
Val	Gln	Val	Leu	His	Arg	Thr	Gly	Leu	Ser	Asn	Pro	Asp	Gly	Leu
3095						3100					3105			
Ala	Val	Asp	Trp	Val	Gly	Gly	Asn	Leu	Tyr	Trp	Cys	Asp	Lys	Gly
3110						3115					3120			
Arg	Asp	Thr	Ile	Glu	Val	Ser	Lys	Leu	Asn	Gly	Ala	Tyr	Arg	Thr
3125						3130					3135			
Val	Leu	Val	Ser	Ser	Gly	Leu	Arg	Glu	Pro	Arg	Ala	Leu	Val	Val
3140						3145					3150			
Asp	Val	Gln	Asn	Gly	Tyr	Leu	Tyr	Trp	Thr	Asp	Trp	Gly	Asp	His
3155						3160					3165			
Ser	Leu	Ile	Gly	Arg	Ile	Gly	Met	Asp	Gly	Ser	Gly	Arg	Ser	Ile
3170						3175					3180			
Ile	Val	Asp	Thr	Lys	Ile	Thr	Trp	Pro	Asn	Gly	Leu	Thr	Val	Asp
3185						3190					3195			
Tyr	Val	Thr	Glu	Arg	Ile	Tyr	Trp	Ala	Asp	Ala	Arg	Glu	Asp	Tyr
3200						3205					3210			
Ile	Glu	Phe	Ala	Ser	Leu	Asp	Gly	Ser	Asn	Arg	His	Val	Val	Leu
3215						3220					3225			
Ser	Gln	Asp	Ile	Pro	His	Ile	Phe	Ala	Leu	Thr	Leu	Phe	Glu	Asp
3230						3235					3240			
Tyr	Val	Tyr	Trp	Thr	Asp	Trp	Glu	Thr	Lys	Ser	Ile	Asn	Arg	Ala
3245						3250					3255			
His	Lys	Thr	Thr	Gly	Ala	Asn	Lys	Thr	Leu	Leu	Ile	Ser	Thr	Leu

3260						3265						3270			
His Arg	Pro Met	Asp Leu	His Val	Phe His	Ala Leu	Arg Gln	Pro								
3275			3280			3285									
Asp Val	Pro Asn	His Pro	Cys Lys	Val Asn	Asn Gly	Gly Cys	Ser								
3290			3295			3300									
Asn Leu	Cys Leu	Leu Ser	Pro Gly	Gly Gly	His Lys	Cys Ala	Cys								
3305			3310			3315									
Pro Thr	Asn Phe	Tyr Leu	Gly Gly	Asp Gly	Arg Thr	Cys Val	Ser								
3320			3325			3330									
Asn Cys	Thr Ala	Ser Gln	Phe Val	Cys Lys	Asn Asp	Lys Cys	Ile								
3335			3340			3345									
Pro Phe	Trp Trp	Lys Cys	Asp Thr	Glu Asp	Asp Cys	Gly Asp	His								
3350			3355			3360									
Ser Asp	Glu Pro	Pro Asp	Cys Pro	Glu Phe	Lys Cys	Arg Pro	Gly								
3365			3370			3375									
Gln Phe	Gln Cys	Ser Thr	Gly Ile	Cys Thr	Asn Pro	Ala Phe	Ile								
3380			3385			3390									
Cys Asp	Gly Asp	Asn Asp	Cys Gln	Asp Asn	Ser Asp	Glu Ala	Asn								
3395			3400			3405									
Cys Asp	Ile His	Val Cys	Leu Pro	Ser Gln	Phe Lys	Cys Thr	Asn								
3410			3415			3420									
Thr Asn	Arg Cys	Ile Pro	Gly Ile	Phe Arg	Cys Asn	Gly Gln	Asp								
3425			3430			3435									
Asn Cys	Gly Asp	Gly Glu	Asp Glu	Arg Asp	Cys Pro	Glu Val	Thr								
3440			3445			3450									
Cys Ala	Pro Asn	Gln Phe	Gln Cys	Ser Ile	Thr Lys	Arg Cys	Ile								
3455			3460			3465									
Pro Arg	Val Trp	Val Cys	Asp Arg	Asp Asn	His Cys	Val Asp	Gly								
3470			3475			3480									
Ser Asp	Glu Pro	Ala Asn	Cys Thr	Gln Met	Thr Cys	Gly Val	Asp								

3485		3490		3495
Glu Phe Arg Cys Lys Asp Ser Gly Arg Cys Ile Pro Ala Arg Trp				
3500		3505		3510
Lys Cys Asp Gly Glu Asp Asp Cys Gly Asp Gly Ser Asp Glu Pro				
3515		3520		3525
Lys Glu Glu Cys Asp Glu Arg Thr Cys Glu Pro Tyr Gln Phe Arg				
3530		3535		3540
Cys Lys Asn Asn Arg Cys Val Pro Gly Arg Trp Gln Cys Asp Tyr				
3545		3550		3555
Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu Glu Ser Cys Thr Pro				
3560		3565		3570
Arg Pro Cys Ser Glu Ser Glu Phe Phe Cys Ala Asn Gly Arg Cys				
3575		3580		3585
Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp Cys Ala Asp				
3590		3595		3600
Gly Ser Asp Glu Lys Asp Cys Thr Pro Arg Cys Asp Met Asp Gln				
3605		3610		3615
Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro Cys				
3620		3625		3630
Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys				
3635		3640		3645
Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn				
3650		3655		3660
Asn Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp				
3665		3670		3675
Asp Cys Gly Asp Asn Ser Asp Glu Asn Pro Glu Glu Cys Ala Arg				
3680		3685		3690
Phe Ile Cys Pro Pro Asn Arg Pro Phe Arg Cys Lys Asn Asp Arg				
3695		3700		3705
Val Cys Leu Trp Ile Gly Arg Gln Cys Asp Gly Val Asp Asn Cys				
3710		3715		3720

Gly	Asp	Gly	Thr	Asp	Glu	Glu	Asp	Cys	Glu	Pro	Pro	Thr	Ala	Gln
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Asn	Pro	His	Cys	Lys	Asp	Lys	Lys	Glu	Phe	Leu	Cys	Arg	Asn	Gln
3740						3745					3750			
Arg	Cys	Leu	Ser	Ser	Ser	Leu	Arg	Cys	Asn	Met	Phe	Asp	Asp	Cys
3755						3760					3765			
Gly	Asp	Gly	Ser	Asp	Glu	Glu	Asp	Cys	Ser	Ile	Asp	Pro	Lys	Leu
3770						3775					3780			
Thr	Ser	Cys	Ala	Thr	Asn	Ala	Ser	Met	Cys	Gly	Asp	Glu	Ala	Arg
3785						3790					3795			
Cys	Val	Arg	Thr	Glu	Lys	Ala	Ala	Tyr	Cys	Ala	Cys	Arg	Ser	Gly
3800						3805					3810			
Phe	His	Thr	Val	Pro	Gly	Gln	Pro	Gly	Cys	Gln	Asp	Ile	Asn	Glu
3815						3820					3825			
Cys	Leu	Arg	Phe	Gly	Thr	Cys	Ser	Gln	Leu	Trp	Asn	Lys	Pro	Lys
3830						3835					3840			
Gly	Gly	His	Leu	Cys	Ser	Cys	Ala	Arg	Asn	Phe	Met	Lys	Thr	His
3845						3850					3855			
Asn	Thr	Cys	Lys	Ala	Glu	Gly	Ser	Glu	Tyr	Gln	Val	Leu	Tyr	Ile
3860						3865					3870			
Ala	Asp	Asp	Asn	Glu	Ile	Arg	Ser	Leu	Phe	Pro	Gly	His	Pro	His
3875						3880					3885			
Ser	Ala	Tyr	Glu	Gln	Thr	Phe	Gln	Gly	Asp	Glu	Ser	Val	Arg	Ile
3890						3895					3900			
Asp	Ala	Met	Asp	Val	His	Val	Lys	Ala	Gly	Arg	Val	Tyr	Trp	Thr
3905						3910					3915			
Asn	Trp	His	Thr	Gly	Thr	Ile	Ser	Tyr	Arg	Ser	Leu	Pro	Pro	Ala
3920						3925					3930			
Ala	Pro	Pro	Thr	Thr	Ser	Asn	Arg	His	Arg	Arg	Gln	Ile	Asp	Arg
3935						3940					3945			

Gly	Val	Thr	His	Leu	Asn	Ile	Ser	Gly	Leu	Lys	Met	Pro	Arg	Gly
3950						3955					3960			
Ile	Ala	Ile	Asp	Trp	Val	Ala	Gly	Asn	Val	Tyr	Trp	Thr	Asp	Ser
3965						3970					3975			
Gly	Arg	Asp	Val	Ile	Glu	Val	Ala	Gln	Met	Lys	Gly	Glu	Asn	Arg
3980						3985					3990			
Lys	Thr	Leu	Ile	Ser	Gly	Met	Ile	Asp	Glu	Pro	His	Ala	Ile	Val
3995						4000					4005			
Val	Asp	Pro	Leu	Arg	Gly	Thr	Met	Tyr	Trp	Ser	Asp	Trp	Gly	Asn
4010						4015					4020			
His	Pro	Lys	Ile	Glu	Thr	Ala	Ala	Met	Asp	Gly	Thr	Leu	Arg	Glu
4025						4030					4035			
Thr	Leu	Val	Gln	Asp	Asn	Ile	Gln	Trp	Pro	Thr	Gly	Leu	Ala	Val
4040						4045					4050			
Asp	Tyr	His	Asn	Glu	Arg	Leu	Tyr	Trp	Ala	Asp	Ala	Lys	Leu	Ser
4055						4060					4065			
Val	Ile	Gly	Ser	Ile	Arg	Leu	Asn	Gly	Thr	Asp	Pro	Ile	Val	Ala
4070						4075					4080			
Ala	Asp	Ser	Lys	Arg	Gly	Leu	Ser	His	Pro	Phe	Ser	Ile	Asp	Val
4085						4090					4095			
Phe	Glu	Asp	Tyr	Ile	Tyr	Gly	Val	Thr	Tyr	Ile	Asn	Asn	Arg	Val
4100						4105					4110			
Phe	Lys	Ile	His	Lys	Phe	Gly	His	Ser	Pro	Leu	Tyr	Asn	Leu	Thr
4115						4120					4125			
Gly	Gly	Leu	Ser	His	Ala	Ser	Asp	Val	Val	Leu	Tyr	His	Gln	His
4130						4135					4140			
Lys	Gln	Pro	Glu	Val	Thr	Asn	Pro	Cys	Asp	Arg	Lys	Lys	Cys	Glu
4145						4150					4155			
Trp	Leu	Cys	Leu	Leu	Ser	Pro	Ser	Gly	Pro	Val	Cys	Thr	Cys	Pro
4160						4165					4170			

Asn Gly 4175	Lys Arg Leu Asp	Asn Gly Thr Cys Val 4180	Pro Val Pro Ser 4185
Pro Thr 4190	Pro Pro Pro Asp	Ala Pro Arg Pro Gly 4195	Thr Cys Thr Leu 4200
Gln Cys 4205	Phe Asn Gly Gly	Ser Cys Phe Leu Asn 4210	Ala Arg Arg Gln 4215
Pro Lys 4220	Cys Arg Cys Gln	Pro Arg Tyr Thr Gly 4225	Asp Lys Cys Glu 4230
Leu Asp 4235	Gln Cys Trp Glu	Tyr Cys His Asn Gly 4240	Gly Thr Cys Ala 4245
Ala Ser 4250	Pro Ser Gly Met	Pro Thr Cys Arg Cys 4255	Pro Thr Gly Phe 4260
Thr Gly 4265	Pro Lys Cys Thr	Ala Gln Val Cys Ala 4270	Gly Tyr Cys Ser 4275
Asn Asn 4280	Ser Thr Cys Thr	Val Asn Gln Gly Asn 4285	Gln Pro Gln Cys 4290
Arg Cys 4295	Leu Pro Gly Phe	Leu Gly Asp Arg Cys 4300	Gln Tyr Arg Gln 4305
Cys Ser 4310	Gly Phe Cys Glu	Asn Phe Gly Thr Cys 4315	Gln Met Ala Ala 4320
Asp Gly 4325	Ser Arg Gln Cys	Arg Cys Thr Val Tyr 4330	Phe Glu Gly Pro 4335
Arg Cys 4340	Glu Val Asn Lys	Cys Ser Arg Cys Leu 4345	Gln Gly Ala Cys 4350
Val Val 4355	Asn Lys Gln Thr	Gly Asp Val Thr Cys 4360	Asn Cys Thr Asp 4365
Gly Arg 4370	Val Ala Pro Ser	Cys Leu Thr Cys Ile 4375	Asp His Cys Ser 4380
Asn Gly 4385	Gly Ser Cys Thr	Met Asn Ser Lys Met 4390	Met Pro Glu Cys 4395
Gln Cys	Pro Pro His Met	Thr Gly Pro Arg Cys	Gln Glu Gln Val



4400                      4405                      4410  
 Val Ser Gln Gln Gln Pro Gly His Met Ala Ser Ile Leu Ile Pro  
 4415                      4420                      4425  
 Leu Leu Leu Leu Leu Leu Leu Leu Val Ala Gly Val Val Phe  
 4430                      4435                      4440  
 Trp Tyr Lys Arg Arg Val Arg Gly Ala Lys Gly Phe Gln His Gln  
 4445                      4450                      4455  
 Arg Met Thr Asn Gly Ala Met Asn Val Glu Ile Gly Asn Pro Thr  
 4460                      4465                      4470  
 Tyr Lys Met Tyr Glu Gly Gly Glu Pro Asp Asp Val Gly Gly Leu  
 4475                      4480                      4485  
 Leu Asp Ala Asp Phe Ala Leu Asp Pro Asp Lys Pro Thr Asn Phe  
 4490                      4495                      4500  
 Thr Asn Pro Val Tyr Ala Thr Leu Tyr Met Gly Gly His Gly Ser  
 4505                      4510                      4515  
 Arg His Ser Leu Ala Ser Thr Asp Glu Lys Arg Glu Leu Leu Gly  
 4520                      4525                      4530  
 Arg Gly Pro Glu Asp Glu Ile Gly Asp Pro Leu Ala  
 4535                      4540                      4545  
  
 <210> 3  
 <211> 1474  
 <212> PRT  
 <213> Homo sapiens  
  
 <400> 3  
  
 Met Gly Lys Asn Lys Leu Leu His Pro Ser Leu Val Leu Leu Leu Leu  
 1                      5                      10                      15  
  
 Val Leu Leu Pro Thr Asp Ala Ser Val Ser Gly Lys Pro Gln Tyr Met  
 20                      25                      30  
  
 Val Leu Val Pro Ser Leu Leu His Thr Glu Thr Thr Glu Lys Gly Cys  
 35                      40                      45  
  
 Val Leu Leu Ser Tyr Leu Asn Glu Thr Val Thr Val Ser Ala Ser Leu  
 50                      55                      60

Glu	Ser	Val	Arg	Gly	Asn	Arg	Ser	Leu	Phe	Thr	Asp	Leu	Glu	Ala	Glu	65	70	75	80
Asn	Asp	Val	Leu	His	Cys	Val	Ala	Phe	Ala	Val	Pro	Lys	Ser	Ser	Ser	85	90	95	
Asn	Glu	Glu	Val	Met	Phe	Leu	Thr	Val	Gln	Val	Lys	Gly	Pro	Thr	Gln	100	105	110	
Glu	Phe	Lys	Lys	Arg	Thr	Thr	Val	Met	Val	Lys	Asn	Glu	Asp	Ser	Leu	115	120	125	
Val	Phe	Val	Gln	Thr	Asp	Lys	Ser	Ile	Tyr	Lys	Pro	Gly	Gln	Thr	Val	130	135	140	
Lys	Phe	Arg	Val	Val	Ser	Met	Asp	Glu	Asn	Phe	His	Pro	Leu	Asn	Glu	145	150	155	160
Leu	Ile	Pro	Leu	Val	Tyr	Ile	Gln	Asp	Pro	Lys	Gly	Asn	Arg	Ile	Ala	165	170	175	
Gln	Trp	Gln	Ser	Phe	Gln	Leu	Glu	Gly	Gly	Leu	Lys	Gln	Phe	Ser	Phe	180	185	190	
Pro	Leu	Ser	Ser	Glu	Pro	Phe	Gln	Gly	Ser	Tyr	Lys	Val	Val	Val	Gln	195	200	205	
Lys	Lys	Ser	Gly	Gly	Arg	Thr	Glu	His	Pro	Phe	Thr	Val	Glu	Glu	Phe	210	215	220	
Val	Leu	Pro	Lys	Phe	Glu	Val	Gln	Val	Thr	Val	Pro	Lys	Ile	Ile	Thr	225	230	235	240
Ile	Leu	Glu	Glu	Glu	Met	Asn	Val	Ser	Val	Cys	Gly	Leu	Tyr	Thr	Tyr	245	250	255	
Gly	Lys	Pro	Val	Pro	Gly	His	Val	Thr	Val	Ser	Ile	Cys	Arg	Lys	Tyr	260	265	270	
Ser	Asp	Ala	Ser	Asp	Cys	His	Gly	Glu	Asp	Ser	Gln	Ala	Phe	Cys	Glu	275	280	285	
Lys	Phe	Ser	Gly	Gln	Leu	Asn	Ser	His	Gly	Cys	Phe	Tyr	Gln	Gln	Val	290	295	300	

Lys Thr Lys Val Phe Gln Leu Lys Arg Lys Glu Tyr Glu Met Lys Leu  
 305 310 315 320

His Thr Glu Ala Gln Ile Gln Glu Glu Gly Thr Val Val Glu Leu Thr  
 325 330 335

Gly Arg Gln Ser Ser Glu Ile Thr Arg Thr Ile Thr Lys Leu Ser Phe  
 340 345 350

Val Lys Val Asp Ser His Phe Arg Gln Gly Ile Pro Phe Phe Gly Gln  
 355 360 365

Val Arg Leu Val Asp Gly Lys Gly Val Pro Ile Pro Asn Lys Val Ile  
 370 375 380

Phe Ile Arg Gly Asn Glu Ala Asn Tyr Tyr Ser Asn Ala Thr Thr Asp  
 385 390 395 400

Glu His Gly Leu Val Gln Phe Ser Ile Asn Thr Thr Asn Val Met Gly  
 405 410 415

Thr Ser Leu Thr Val Arg Val Asn Tyr Lys Asp Arg Ser Pro Cys Tyr  
 420 425 430

Gly Tyr Gln Trp Val Ser Glu Glu His Glu Glu Ala His His Thr Ala  
 435 440 445

Tyr Leu Val Phe Ser Pro Ser Lys Ser Phe Val His Leu Glu Pro Met  
 450 455 460

Ser His Glu Leu Pro Cys Gly His Thr Gln Thr Val Gln Ala His Tyr  
 465 470 475 480

Ile Leu Asn Gly Gly Thr Leu Leu Gly Leu Lys Lys Leu Ser Phe Tyr  
 485 490 495

Tyr Leu Ile Met Ala Lys Gly Gly Ile Val Arg Thr Gly Thr His Gly  
 500 505 510

Leu Leu Val Lys Gln Glu Asp Met Lys Gly His Phe Ser Ile Ser Ile  
 515 520 525

Pro Val Lys Ser Asp Ile Ala Pro Val Ala Arg Leu Leu Ile Tyr Ala  
 530 535 540

Val Leu Pro Thr Gly Asp Val Ile Gly Asp Ser Ala Lys Tyr Asp Val

545		550		555		560
Glu Asn Cys Leu Ala Asn Lys Val Asp Leu Ser Phe Ser Pro Ser Gln						
	565			570		575
Ser Leu Pro Ala Ser His Ala His Leu Arg Val Thr Ala Ala Pro Gln						
	580			585		590
Ser Val Cys Ala Leu Arg Ala Val Asp Gln Ser Val Leu Leu Met Lys						
	595			600		605
Pro Asp Ala Glu Leu Ser Ala Ser Ser Val Tyr Asn Leu Leu Pro Glu						
	610			615		620
Lys Asp Leu Thr Gly Phe Pro Gly Pro Leu Asn Asp Gln Asp Asp Glu						
	625			630		635
Asp Cys Ile Asn Arg His Asn Val Tyr Ile Asn Gly Ile Thr Tyr Thr						
		645		650		655
Pro Val Ser Ser Thr Asn Glu Lys Asp Met Tyr Ser Phe Leu Glu Asp						
	660			665		670
Met Gly Leu Lys Ala Phe Thr Asn Ser Lys Ile Arg Lys Pro Lys Met						
	675			680		685
Cys Pro Gln Leu Gln Gln Tyr Glu Met His Gly Pro Glu Gly Leu Arg						
	690			695		700
Val Gly Phe Tyr Glu Ser Asp Val Met Gly Arg Gly His Ala Arg Leu						
	705			710		715
Val His Val Glu Glu Pro His Thr Glu Thr Val Arg Lys Tyr Phe Pro						
		725		730		735
Glu Thr Trp Ile Trp Asp Leu Val Val Val Asn Ser Ala Gly Val Ala						
	740			745		750
Glu Val Gly Val Thr Val Pro Asp Thr Ile Thr Glu Trp Lys Ala Gly						
	755			760		765
Ala Phe Cys Leu Ser Glu Asp Ala Gly Leu Gly Ile Ser Ser Thr Ala						
	770			775		780
Ser Leu Arg Ala Phe Gln Pro Phe Phe Val Glu Leu Thr Met Pro Tyr						
	785			790		795
						800

Ser Val Ile Arg Gly Glu Ala Phe Thr Leu Lys Ala Thr Val Leu Asn  
 805 810 815

Tyr Leu Pro Lys Cys Ile Arg Val Ser Val Gln Leu Glu Ala Ser Pro  
 820 825 830

Ala Phe Leu Ala Val Pro Val Glu Lys Glu Gln Ala Pro His Cys Ile  
 835 840 845

Cys Ala Asn Gly Arg Gln Thr Val Ser Trp Ala Val Thr Pro Lys Ser  
 850 855 860

Leu Gly Asn Val Asn Phe Thr Val Ser Ala Glu Ala Leu Glu Ser Gln  
 865 870 875 880

Glu Leu Cys Gly Thr Glu Val Pro Ser Val Pro Glu His Gly Arg Lys  
 885 890 895

Asp Thr Val Ile Lys Pro Leu Leu Val Glu Pro Glu Gly Leu Glu Lys  
 900 905 910

Glu Thr Thr Phe Asn Ser Leu Leu Cys Pro Ser Gly Gly Glu Val Ser  
 915 920 925

Glu Glu Leu Ser Leu Lys Leu Pro Pro Asn Val Val Glu Glu Ser Ala  
 930 935 940

Arg Ala Ser Val Ser Val Leu Gly Asp Ile Leu Gly Ser Ala Met Gln  
 945 950 955 960

Asn Thr Gln Asn Leu Leu Gln Met Pro Tyr Gly Cys Gly Glu Gln Asn  
 965 970 975

Met Val Leu Phe Ala Pro Asn Ile Tyr Val Leu Asp Tyr Leu Asn Glu  
 980 985 990

Thr Gln Gln Leu Thr Pro Glu Val Lys Ser Lys Ala Ile Gly Tyr Leu  
 995 1000 1005

Asn Thr Gly Tyr Gln Arg Gln Leu Asn Tyr Lys His Tyr Asp Gly  
 1010 1015 1020

Ser Tyr Ser Thr Phe Gly Glu Arg Tyr Gly Arg Asn Gln Gly Asn  
 1025 1030 1035

Thr	Trp	Leu	Thr	Ala	Phe	Val	Leu	Lys	Thr	Phe	Ala	Gln	Ala	Arg
1040						1045					1050			
Ala	Tyr	Ile	Phe	Ile	Asp	Glu	Ala	His	Ile	Thr	Gln	Ala	Leu	Ile
1055						1060					1065			
Trp	Leu	Ser	Gln	Arg	Gln	Lys	Asp	Asn	Gly	Cys	Phe	Arg	Ser	Ser
1070						1075					1080			
Gly	Ser	Leu	Leu	Asn	Asn	Ala	Ile	Lys	Gly	Gly	Val	Glu	Asp	Glu
1085						1090					1095			
Val	Thr	Leu	Ser	Ala	Tyr	Ile	Thr	Ile	Ala	Leu	Leu	Glu	Ile	Pro
1100						1105					1110			
Leu	Thr	Val	Thr	His	Pro	Val	Val	Arg	Asn	Ala	Leu	Phe	Cys	Leu
1115						1120					1125			
Glu	Ser	Ala	Trp	Lys	Thr	Ala	Gln	Glu	Gly	Asp	His	Gly	Ser	His
1130						1135					1140			
Val	Tyr	Thr	Lys	Ala	Leu	Leu	Ala	Tyr	Ala	Phe	Ala	Leu	Ala	Gly
1145						1150					1155			
Asn	Gln	Asp	Lys	Arg	Lys	Glu	Val	Leu	Lys	Ser	Leu	Asn	Glu	Glu
1160						1165					1170			
Ala	Val	Lys	Lys	Asp	Asn	Ser	Val	His	Trp	Glu	Arg	Pro	Gln	Lys
1175						1180					1185			
Pro	Lys	Ala	Pro	Val	Gly	His	Phe	Tyr	Glu	Pro	Gln	Ala	Pro	Ser
1190						1195					1200			
Ala	Glu	Val	Glu	Met	Thr	Ser	Tyr	Val	Leu	Leu	Ala	Tyr	Leu	Thr
1205						1210					1215			
Ala	Gln	Pro	Ala	Pro	Thr	Ser	Glu	Asp	Leu	Thr	Ser	Ala	Thr	Asn
1220						1225					1230			
Ile	Val	Lys	Trp	Ile	Thr	Lys	Gln	Gln	Asn	Ala	Gln	Gly	Gly	Phe
1235						1240					1245			
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